IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re patent application of **David S. TERMAN**

Serial No.: 09/751/708

Filed: December 28, 2000

MAR 0.1 2002 SS

Group Art Unit: 1635

Examiner:

For: COMPOSITIONS AND METHODS FOR TREATMENT OF NEOPLASTIC DISEASE

STATEMENT IN ACCORDANCE WITH 37 CFR 1.821 (f)

Assistant Commissioner for Patents Washington, D.C. 20231
BOX SEQUENCE

Sir:

I hereby state that the information recorded in computer readable form is identical to the written (paper) sequence listing.

Respectfully submitted,

'Clean

2/26/02 Date

David S. Terman

SEQUENCE LISTING <110> TERMAN, David S <120> COMPOSITIONS AND METHODS FOR TREATMENT OF NEOPLASTIC DISEASE <130> 870759 <140> US 09/870,759 <141> 2001-05-30 <150> US 60/208,128 <170> PatentIn version 3.1 <210> 1 <151> 2000-05-30 <160> 166 <211> 801 <212> DNA <213> Staphylococcus sp. <400> 1 atgtataaga gattatttat ttcacatgta attttgatat tcgcactgat attagttatt 60 120 totacaccca acgttttago agagagtcaa coagatocta aaccagatga gttgcacaaa tcgagtaaat tcactggttt gatggaaaat atgaaagttt tgtatgatga taatcatgta 180 tcagcaataa acgttaaatc tatagatcaa tttctatact ttgacttaat atattctatt 240 300 aaggacacta agttagggaa ttatgataat gttcgagtcg aatttaaaaa caaagattta qctgataaat acaaagataa atacgtagat gtgtttggag ctaattatta ttatcaatgt 360 tatttttcta aaaaaacqaa tgatattaat tcgcatcaaa ctgacaaacg aaaaacttgt 420 atgtatggtg gtgtaactga gcataatgga aaccaattag ataaatatag aagtattact 480 540 gttcgggtat ttgaagatgg taaaaattta ttatcttttg acgtacaaac taataagaaa aaggtgactg ctcaagaatt agattaccta actcgtcact atttggtgaa aaataaaaaa 600 ctctatgaat ttaacaactc gccttatgaa acgggatata ttaaaatttat agaaaatgag 660 aataqctttt qqtatqacat qatqcctqca ccaqqaqata aatttgacca atctaaatat 720 ttaatgatgt acaatgacaa taaaatggtt gattctaaag atgtgaagat tgaagtttat 780 801 cttacgacaa agaaaaagtg a

MAR 0 1 2002

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gct Ala 225	caa Gln	ggg	caa Gln	tat Tyr	cca Pro 230	gat Asp	aca Thr	tta Leu	tta Leu	aga Arg 235	att Ile	tac Tyr	aga Arg	gat Asp	aat Asn 240	720
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1	-	Буз		5					10					15		
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Lys Ile Tyr Ile Phe Phe Met Arg Val Thr Leu Val Thr His Glu Asn 50 55 60

Val Lys Ser Val Asp Gln Leu Leu Ser His Asp Leu Ile Tyr Asn Val 65 70 75 80

Ser Gly Pro Asn Tyr Asp Lys Leu Lys Thr Glu Leu Lys Asn Gln Glu 85 90 95

Met Ala Thr Leu Phe Lys Asp Lys Asn Val Asp Ile Tyr Gly Val Glu
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Tyr Tyr His Leu Cys Tyr Leu Cys Glu Asn Ala Glu Arg Ser Ala Cys 115 120 125

Leu Tyr Gly Gly Val Thr Asn His Glu Gly Asn His Leu Glu Ile Pro 130 135 140

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Phe Asp Ile Glu Gln Ile Lys Asn Gly Asn Cys Ser Arg Ile Ser Tyr 165 170 175

Thr Val Arg Lys Tyr Leu Thr Asp Asn Lys Gln Leu Tyr Thr Asn Gly 180 185

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Glu Ser Phe Trp Phe Asp Phe Phe Pro Glu Pro Glu Phe Thr Gln Ser 210 215 220

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Leu Glu Arg Ser Gly Met Gln Ile Ser Arg His Ser Leu Val Ser Ser 145 150 155 160

Tyr Leu Ala Leu Met Glu Phe Ser Gly Asn Thr Met Thr Arg Asp Ala 165 170 175

Ser Arg Ala Val Leu Arg Phe Val Thr Val Thr Ala Glü Ala Leu Arg 180 185 190

Phe Arg Gln Ile Gln Arg Glu Phe Arg Gln Ala Leu Ser Glu Thr Ala 195 200 205

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Arg Val Gly Arg Ile Ser Phe Asn Asn Ile Ser Ala Ile Leu Gly Thr 245 250 255

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	_	-	_	_	aat Asn				_						_	1244	
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Ala Ser Glu Lys Tyr Asp Ala Asn Ala Pro Gln Gln Val Gln Pro Tyr 50 55 60

Ser Val Ser Ser Ser Ala Phe Glu Asn Leu His Pro Asn Asn Glu Met 65 70 75 80

Glu Ser Ser Ile Asn Pro Phe Ser Ala Ser Asp Thr Glu Arg Asn Ala 85 90 95

Ala Ile Ile Asp Arg Ala Asn Lys Glu Gln Glu Thr Glu Ala Val Asn 100 105 110

Lys Met Ile Ser Thr Gly Ala Arg Leu Ala Ala Ser Gly Arg Ala Ser 115 120 125

Asp Val Ala His Ser Met Val Gly Asp Ala Val Asn Gln Glu Ile Lys 130 135 140

Gln Trp Leu Asn Arg Phe Gly Thr Ala Gln Val Asn Leu Asn Phe Asp 145 150 155 160

Lys Asn Phe Ser Leu Lys Glu Ser Ser Leu Asp Trp Leu Ala Pro Trp 165 170 175

Tyr Asp Ser Ala Ser Phe Leu Phe Phe Ser Gln Leu Gly Ile Arg Asn 180 185 190

Lys Asp Ser Arg Asn Thr Leu Asn Leu Gly Val Gly Ile Arg Thr Leu 195 200 205

Glu Asn Gly Trp Leu Tyr Gly Leu Asn Thr Phe Tyr Asp Asn Asp Leu 210 215 220

- Thr Gly His Asn His Arg Ile Gly Leu Gly Ala Glu Ala Trp Thr Asp 225 230 235 235
- Tyr Leu Gln Leu Ala Ala Asn Gly Tyr Phe Arg Leu Asn Gly Trp His 245 250 255
- Ser Ser Arg Asp Phe Ser Asp Tyr Lys Glu Arg Pro Ala Thr Gly Gly 260 265 270
- Asp Leu Arg Ala Asn Ala Tyr Leu Pro Ala Leu Pro Gln Leu Gly Gly 275 280 285
- Lys Leu Met Tyr Glu Gln Tyr Thr Gly Glu Arg Val Ala Leu Phe Gly 290 295 300
- Lys Asp Asn Leu Gln Arg Asn Pro Tyr Ala Val Thr Ala Gly Ile Asn 305 310 315 320
- Tyr Thr Pro Val Pro Leu Leu Thr Val Gly Val Asp Gln Arg Met Gly 325 330 335
- Lys Ser Ser Lys His Glu Thr Gln Trp Asn Leu Gln Met Asn Tyr Arg 340 345 350
- Leu Gly Glu Ser Phe Gln Ser Gln Leu Ser Pro Ser Ala Val Ala Gly 355 360 365
- Thr Arg Leu Leu Ala Glu Ser Arg Tyr Asn Leu Val Asp Arg Asn Asn 370 375 380
- Asn Ile Val Leu Glu Tyr Gln Lys Gln Gln Val Val Lys Leu Thr Leu 385 390 395 400
- Ala Gl
n Val Gl
n Gly Ala Ser Ala Val Arg Glu Ile Val Tr
p Ser Asp $420 \hspace{1.5cm} 425 \hspace{1.5cm} 430 \hspace{1.5cm}$
- Ala Glu Leu Ile Ala Ala Gly Gly Thr Leu Thr Pro Leu Ser Thr Thr 435 440 445
- Gln Phe Asn Leu Val Leu Pro Pro Tyr Lys Arg Thr Ala Gln Val Ser 450 455 460

Arg Val Thr Asp Asp Leu Thr Ala Asn Phe Tyr Ser Leu Ser Ala Leu 465 470 475 480

Ala Val Asp His Gln Gly Asn Arg Ser Asn Ser Phe Thr Leu Ser Val 485 490 495

Thr Val Gln Gln Pro Gln Leu Thr Leu Thr Ala Ala Val Ile Gly Asp 500 505 510

Gly Ala Pro Ala Asn Gly Lys Thr Ala Ile Thr Val Glu Phe Thr Val 515 525

Ala Asp Phe Glu Gly Lys Pro Leu Ala Gly Gln Glu Val Val Ile Thr 530 535 540

Thr Asn Asn Gly Ala Leu Pro Asn Lys Ile Thr Glu Lys Thr Asp Ala 545 550 555 560

Asn Gly Val Ala Arg Ile Ala Leu Thr Asn Thr Thr Asp Gly Val Thr 565 570 575

Val Val Thr Ala Glu Val Glu Gly Gln Arg Gln Ser Val Asp Thr His 580 585 590

Phe Val Lys Gly Thr Ile Ala Ala Asp Lys Ser Thr Leu Ala Ala Val595 \cdot 600 605

Pro Thr Ser Ile Ile Ala Asp Gly Leu Met Ala Ser Thr Ile Thr Leu 610 615 620

Glu Leu Lys Asp Thr Tyr Gly Asp Pro Gln Ala Gly Ala Asn Val Ala 625 630 635 640

Phe Asp Thr Thr Leu Gly Asn Met Gly Val Ile Thr Asp His Asn Asp 645 650 655

Gly Thr Tyr Ser Ala Pro Leu Thr Ser Thr Thr Leu Gly Val Ala Thr 660 665 670

Val Thr Val Lys Val Asp Gly Ala Ala Phe Ser Val Pro Ser Val Thr 675 680 685

Val Asn Phe Thr Ala Asp Pro Ile Pro Asp Ala Gly Arg Ser Ser Phe 690 695 700

Thr Val Ser Thr Pro Asp Ile Leu Ala Asp Gly Thr Met Ser Ser Thr 705 710 715 720

Leu Ser Phe Val Pro Val Asp Lys Asn Gly His Phe Ile Ser Gly Met 725 730 735

Gln Gly Leu Ser Phe Thr Gln Asn Gly Val Pro Val Ser Ile Ser Pro 740 745 750

Ile Thr Glu Gln Pro Asp Ser Tyr Thr Ala Thr Val Val Gly Asn Ser $755 \hspace{1.5cm} 760 \hspace{1.5cm} 765$

Val Gly Asp Val Thr Ile Thr Pro Gln Val Asp Thr Leu Ile Leu Ser 770 780

Thr Leu Gln Lys Lys Ile Ser Leu Phe Pro Val Pro Thr Leu Thr Gly 785 790 795 800

Ile Leu Val As
n Gly Gl
n As
n Phe Ala Thr Asp Lys Gly Phe Pro Lys 805 810 815

Thr Ile Phe Lys Asn Ala Thr Phe Gln Leu Gln Met Asp Asn Asp Val 820 825 830

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Ser Val Asn Asp Gln Gly Gln Val Thr Ile Thr Tyr Gln Thr Tyr Ser 850 860

Glu Val Ala Val Thr Ala Lys Ser Lys Lys Phe Pro Ser Tyr Ser Val 865 870 875 880

Ser Tyr Arg Phe Tyr Pro Asn Arg Trp Ile Tyr Asp Gly Gly Arg Ser 885 890 895

Leu Val Ser Ser Leu Glu Ala Ser Arg Gln Cys Gln Gly Ser Asp Met 900 905 910

Ser Ala Val Leu Glu Ser Ser Arg Ala Thr Asn Gly Thr Arg Ala Pro 915 920 925

Asp Gly Thr Leu Trp Gly Glu Trp Gly Ser Leu Thr Ala Tyr Ser Ser 930 935 940

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tcggcaaagc agccggatac cgctacgtat cttgaagtat tgaaaatatt acgatgcaaa 60 aaagaaaatt taagtataat acagcaggat tctttaacgg attcttaaca attttctaa 120 ctgaccataa aggaaccaaa at atg aaa aaa gca ctt gcc aca ctg att gcc Met Lys Lys Ala Leu Ala Thr Leu Ile Ala 1 5 10 ctc gct ctc ccg gcc gcc gca ctg gcg gaa ggc gca tcc ggc ttt tac Leu Ala Leu Pro Ala Ala Ala Leu Ala Glu Gly Ala Ser Gly Phe Tyr 15 20 25 gtc caa gcc gat gcc gca cac gca aaa gcc tca agc tct tta ggt tct Val Gln Ala Asp Ala Ala His Ala Lys Ala Ser Ser Ser Leu Gly Ser
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- Phe Leu Thr Gly Ile Val Gly Asn Gly Leu Val Ile Leu Val Met Gly 50 60
- Tyr Gln Lys Lys Leu Arg Ser Met Thr Asp Lys Tyr Arg Leu His Leu 65 75 80
- Ser Val Ala Asp Leu Leu Phe Val Ile Thr Leu Pro Phe Trp Ala Val 85 90 95
- Asp Ala Val Ala Asn Trp Tyr Phe Gly Asn Phe Leu Cys Lys Ala Val
 100 105 110
- His Val Ile Tyr Thr Val Asn Leu Tyr Ser Ser Val Leu Ile Leu Ala 115 120 125
- Phe Ile Ser Leu Asp Arg Tyr Leu Ala Ile Val His Ala Thr Asn Ser 130 135 140
- Gln Arg Pro Arg Lys Leu Leu Ala Glu Lys Val Val Tyr Val Gly Val 145 150 155 160

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ctg Leu	gtg Val	acc Thr 435	agt Ser	Gly	gct Ala	gtä Val	agc Ser 440	agt Ser	ggg Gly	aac Asn	ctt Leu	gtc Val 445	aca Thr	ttt Phe	gtt Val	1344
ctc Leu	tac Tyr 450	cag Gln	atg Met	cag Gln	ttc Phe	acc Thr 455	cag Gln	gct Ala	ttg Leu	gag Glu	gta Val 460	ctg Leu	ctc Leu	tcc Ser	atc Ile	1392
tac Tyr 465	Pro	aga Arg	gta Val	cag Gln	aag Lys 470	gct Ala	gtg Val	ggc Gly	tcc Ser	tca Ser 475	gag Glu	aaa Lys	ata Ile	ttt Phe	gag Glu 480	1440
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					ctg Leu 550											1680
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cct Pro	gag Glu 690	cgg Arg	tac Tyr	tcc Ser	cgc Arg	tca Ser 695	gtg Val	ctt Leu	çtc Leu	atc Ile	acc Thr 700	cag Gln	cac His	ctc Leu	agc Ser	2112
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					His					Glu						2208
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Gly Ser Lys Ser Glu Asn Ala Gly Ala Gln Gly Trp Leu Ala Ala Leu 85 90 95

Lys Pro Leu Ala Ala Ala Leu Gly Leu Ala Leu Pro Gly Leu Ala Leu 100 105 110

Phe Arg Glu Leu Ile Ser Trp Gly Ala Pro Gly Ser Ala Asp Ser Thr 115 120 125

Arg Leu Leu His Trp Gly Ser His Pro Thr Ala Phe Val Val Ser Tyr 130 135 140

Ala Ala Leu Pro Ala Ala Ala Leu Trp His Lys Leu Gly Ser Leu 145 150 155 160

Trp Val Pro Gly Gly Gln Gly Gly Ser Gly Asn Pro Val Arg Arg Leu 165 170 175

Leu Gly Cys Leu Gly Ser Glu Thr Arg Arg Leu Ser Leu Phe Leu Val 180 185 190

Leu Val Val Leu Ser Ser Leu Gly Glu Met Ala Ile Pro Phe Thr 195 200 205

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Thr Arg Asn Leu Thr Leu Met Ser Ile Leu Thr Ile Ala Ser Ala Val 225 230 235 240

Leu Glu Phe Val Gly Asp Gly Ile Tyr Asn Asn Thr Met Gly His Val 245 250 255

His Ser His Leu Gln Gly Glu Val Phe Gly Ala Val Leu Arg Gln Glu 265 270

Thr Glu Phe Phe Gln Gln Asn Gln Thr Gly Asn Ile Met Ser Arg Val 275 280 285

Thr Glu Asp Thr Ser Thr Leu Ser Asp Ser Leu Ser Glu Asn Leu Ser 290 295 300

Leu Phe Leu Trp Tyr Leu Val Arg Gly Leu Cys Leu Leu Gly Ile Met 305 310 315 320

Leu Trp Gly Ser Val Ser Leu Thr Met Val Thr Leu Val Thr Leu Pro 325 330 335

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Glu Val Gln Val Arg Glu Ser Leu Ala Lys Ser Ser Gln Val Ala Ile 355 360 365

Glu Ala Leu Ser Ala Met Pro Thr Val Arg Ser Phe Ala Asn Glu Glu 370 375 380

Gly Glu Ala Gln Lys Phe Arg Glu Lys Leu Gln Glu Ile Lys Thr Leu 385 390 395 400

Asn Gln Lys Glu Ala Val Ala Tyr Ala Val Asn Ser Trp Thr Thr Ser 405 410 415

Ile Ser Gly Met Leu Leu Lys Val Gly Ile Leu Tyr Ile Gly Gln 420 425 430

Leu Val Thr Ser Gly Ala Val Ser Ser Gly Asn Leu Val Thr Phe Val 435 440 445

Leu Tyr Gln Met Gln Phe Thr Gln Ala Leu Glu Val Leu Leu Ser Ile 450 455 460 Tyr Pro Arg Val Gln Lys Ala Val Gly Ser Ser Glu Lys Ile Phe Glu 465 470 475 480

Tyr Leu Asp Arg Thr Pro Arg Cys Pro Pro Ser Gly Leu Leu Thr Pro 485 490 495

Leu His Leu Glu Gly Leu Val Gln Phe Gln Asp Val Ser Phe Ala Tyr 500 505 510

Pro Asn Arg Pro Asp Val Leu Val Leu Gln Gly Leu Thr Phe Thr Leu 515 520 525

Arg Pro Gly Glu Val Thr Ala Leu Val Gly Pro Asn Gly Ser Gly Lys 530 535 540

Ser Thr Val Ala Ala Leu Leu Gln Asn Leu Tyr Gln Pro Thr Gly Gly 545 550 555 560

Gln Leu Leu Leu Asp Gly Lys Pro Leu Pro Gln Tyr Glu His Arg Tyr 565 570 575

Leu His Arg Gln Val Ala Ala Val Gly Gln Glu Pro Gln Val Phe Gly 580 585 590

Arg Ser Leu Gln Glu Asn Ile Ala Tyr Gly Leu Thr Gln Lys Pro Thr 595 600 605

Met Glu Glu Ile Thr Ala Ala Ala Val Lys Ser Gly Ala His Ser Phe 610 615 620

Ile Ser Gly Leu Pro Gln Gly Tyr Asp Thr Glu Val Gly Glu Ala Gly 625 630 635 640

Ser Gln Leu Ser Gly Gly Gln Gln Gln Ala Val Ala Leu Ala Arg Ala 645 650 655

Leu Ilé Arg Lys Pro Cys Val Leu Ile Leu Asp Asp Ala Thr Ser Ala 660 665 670

Leu Asp Ala Asn Ser Gln Leu Gln Val Glu Gln Leu Leu Tyr Glu Ser 675 680 685

Pro Glu Arg Tyr Ser Arg Ser Val Leu Leu Ile Thr Gln His Leu Ser 690 695 700

Leu Val Glu Gln Ala Asp His Ile Leu Phe Leu Glu Gly Gly Ala Ile 705 710 715 720	
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cat His 240	ttt Phe	att Ile	gct Ala	gag Glu	ttt Phe 245	aag Lys	cgc Arg	aag Lys	cat His	aag Lys 250	aag Lys	gac Asp	atc Ile	agt Ser	gag Glu 255	829
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2085

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Gln Gly Asn Arg Thr Thr Pro Ser Tyr Val Ala Phe Thr Asp Thr Glu
35 40 45

Arg Leu Ile Gly Asp Ala Ala Lys Asn Gln Val Ala Met Asn Pro Thr 50 55 60

Asn Thr Val Phe Asp Ala Lys Arg Leu Ile Gly Arg Arg Phe Asp Asp 65 70 75 80

Ala Val Val Gln Ser Asp Met Lys His Trp Pro Phe Met Val Val Asn 85 90 95

Asp Ala Gly Arg Pro Lys Val Gln Val Glu Tyr Lys Gly Glu Thr Lys 100 105 110

Ser Phe Tyr Pro Glu Glu Val Ser Ser Met Val Leu Thr Lys Met Lys 115 120 125

Glu Ile Ala Glu Ala Tyr Leu Gly Lys Thr Val Thr Asn Ala Val Val 130 \$135\$

Thr Val Pro Ala Tyr Phe Asn Asp Ser Gln Arg Gln Ala Thr Lys Asp 145 150 155 160

Ala Gly Thr Ile Ala Gly Leu Asn Val Leu Arg Ile Ile Asn Glu Pro 165 170 175

Thr Ala Ala Ile Ala Tyr Gly Leu Asp Lys Lys Val Gly Ala Glu 180 185 190

Arg Asn Val Leu Ile Phe Asp Léu Gly Gly Gly Thr Phe Asp Val Ser

Ile Leu Thr Ile Glu Asp Gly Ile Phe Glu Val Lys Ser Thr Ala Gly

Asp Thr His Leu Gly Gly Glu Asp Phe Asp Asn Arg Met Val Asn His 225 230 235 240

220

Phe Ile Ala Glu Phe Lys Arg Lys His Lys Lys Asp Ile Ser Glu Asn 245 250 255

Lys Arg Ala Val Arg Arg Leu Arg Thr Ala Cys Glu Arg Ala Lys Arg 260 265 270

Thr Leu Ser Ser Ser Thr Gln Ala Ser Ile Glu Ile Asp Ser Leu Tyr 275 280 285

Glu Gly Ile Asp Phe Tyr Thr Ser Ile Thr Arg Ala Arg Phe Glu Glu 290 295 300

Leu Asn Ala Asp Leu Phe Arg Gly Thr Leu Asp Pro Val Glu Lys Ala 305 310 315 320

Val Gly Gly Ser Thr Arg Ile Pro Lys Ile Gln Lys Leu Leu Gln Asp $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$

Phe Phe Asn Gly Lys Glu Leu Asn Lys Ser Ile Asn Pro Asp Glu Ala 355 . 360 365

Val Ala Tyr Gly Ala Ala Val Gln Ala Ala Ile Leu Ser Gly Asp Lys 370 375 380

Ser Glu Asn Val Gln Asp Leu Leu Leu Leu Asp Val Thr Pro Leu Ser 385 390 395 400

Leu Gly Ile Glu Thr Ala Gly Gly Val Met Thr Val Leu Ile Lys Arg \$405\$

Asn Thr Thr Ile Pro Thr Lys Gln Thr Gln Thr Phe Thr Thr Tyr Ser

Asp Asn Gln Pro Gly Val Leu Ile Gln Val Tyr Glu Gly Glu Arg Ala 435 440 445

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Glu	Asp	Ile 515	Glu	Arg	Met	Val	Gln 520	Glu	Ala	Glu	Lys	Tyr 525	Lys	Ala	Glu	
Asp	Glu 530	Lys	Gln	Arg	Asp	L ys 535	Val	Ser	Ser	Lys	Asn 540	Ser	Leu	Glu	Ser	
Tyr 545	Ala	Phe	Asn	Met	Lys 550	Ala	Thr	Val	Glu	Asp 555	Glu	Lys	Leu	Gln	Gly 560	
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Met Asn Lys Asn Val Leu Lys Phe 1

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				Asn	caa Gln											1566
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Lys	Lys	Leu 475	Glų	Asp	Gly	Thr	Thr 480	Lys	Val	Thr	Trp	Ser 485	Asn	Leu	Pro	
gaa Glu	aat Asn 490	gac Asp	aaa Lys	aat Asn	ggc Gly	aag Lys 495	gct Ala	att Ile	aaa Lys	tat Tyr	tta Leu 500	gtt Val	aaa Lys	gaa Glu	gta Val	1662
aat Asn 505	gct Ala	caa Gln	ggt Gly	gaa Glu	gat Asp 510	aca Thr	aca Thr	cca Pro	gaa Glu	gga Gly 515	tat Tyr	act Thr	aaa Lys	aaa Lys	gaa Glu 520	1710
aat Asn	ggt Gly	tta Leu	gtg Val	gtt Val 525	act Thr	aat Asn	act Thr	gaa Glu	aaa Lys 530	cca Pro	atc Ile	gaa Glu	aca Thr	aca Thr 535	tca Ser	1758
					gta Val											1806
aga Arg	cca Pro	gaa Glu 555	aaa Lys	gtc Val	agt Ser	gtg Val	aat Asn 560	tta Leu	ttg Leu	ġct Ala	aac Asn	ggg Gly 565	gag Glu	aaa Lys	gta Val	1854
aaa Lys	acg Thr 570	tta Lėu	gac Asp	gtg Val	aca Thr	tct Ser 575	gaa Glu	aca Thr	aac Asn	tgg Trp	aag Lys 580	tac Tyr	gaa Glu	ttt Phe	aaa Lys	1902
					gat Asp 590											1950
_	_		_		gac Asp				-				_			1998
					cca Pro											2046
tgg Trp	gat Asp	gac Asp 635	aat Asn	aat Asn	aac Asn	caa Gln	gac Asp 640	gga Gly	aaa Lys	cga Arg	cca Pro	act Thr 645	gaa Glu	atc Ile	aaa Lys	2094
gtt Val	gag Glu 650	tta Leu	tat Tyr	caa Gln	gat Asp	gga Gly 655	aaa Lys	gca Ala	aca Thr	gga Gly	aaa Lys 660	acg Thr	gca Ala	ata Ile	tta Leu	2142
aat Asn 665	gaa Glu	tct Ser	aat Asn	aac Asn	tgg Trp 670	aca Thr	cat His	acg Thr	tgg Trp	aca Thr 675	gga Gly	tta Leu	gat Asp	gaa Glu	aaa Lys 680	2190
gca Ala	aaa Lys	gga Gly	caa Gln	caa Gln 685	gta Val	aaa Lys	tac Tyr	aca Thr	gtc Val 690	gag Glu	gaa Glu	tta Leu	aca Thr	aag Lys 695	gtc Val	2238
					cat His											2286
gtg	acg	aat	aaa	tat	acg	cca	gaa	aca	aca	tca	att	agt	ggt	gaa	aaa	2334

Val	Thr	Asn 715	Lys	Tyr	Thr	Pro	Glu 720	Thr	Thr	Ser	Ile	Ser 725	Gly	Glu	Lys	
gta Val	tgg Trp 730	gac Asp	gac Asp	aaa Lys	gac Asp	aat Asn 735	caa Gln	gat Asp	ggt Gly	aag Lys	aga Arg 740	cca Pro	gaa Glu	aaa Lys	gtc Val	2382
agt Ser 745	gtg Val	aat Asn	tta Leu	ttg Leu	gct Ala 750	gat Asp	gga Gly	gag Glu	aaa Lys	gta Val 755	aaa Lys	acg Thr	tta Leu	gac Asp	gtg Val 760	2430
aca Thr	tct Ser	gaa Glu	aca Thr	aac Asn 765	tgg Trp	aag Lys	tac Tyr	gaa Glu	ttt Phe 770	aaa Lys	gac Asp	tta Leu	ccg Pro	aag Lys 775	tat Tyr	2478
gat Asp	gaa Glu	gga Gly	aag Lys 780	aaa Lys	ata Ile	gaa Glu	tat Tyr	aca Thr 785	gtg Val	acc Thr	gaa Glu	gat Asp	cac His 790	gta Val	aaa Lys	2526
gac Asp	tac Tyr	aca Thr 795	aca Thr	gac Asp	atc Ile	aac Asn	ggt Gly 800	acg Thr	aca Thr	ata Ile	acg Thr	aac Asn 805	aag Lys	tat Tyr	aca Thr	2574
cca Pro	gga Gly 810	gag Glu	aca Thr	tcg Ser	gca Ala	aca Thr 815	gta Val	aca Thr	aaa Lys	aat Asn	tgg Trp 820	gat Asp	gac Asp	aat Asn	aat Asn	2622
aac Asn 825	caa Gln	gac Asp	gga Gly	aaa Lys	cga Arg 830	cca Pro	act Thr	gaa Glu	atc Ile	aaa Lys 835	gtt Val	gag Glu	tta Leu	tat Tyr	caa Gln 840	2670
gat Asp	gga Gly	aaa Lys	gċa Ala	aca Thr 845	gga Gly	aaa Lys	acg Thr	gca Ala	ata Ile 850	tta Leu	aat Asn	gaa Glu	tct Ser	aat Asn 855	aac Asn	2718
tgg Trp	aca Thr	cat His	acg Thr 860	tgg Trp	aca Thr	gga Gly	tta Leu	gat Asp 865	gaa Glu	aaa Lys	gca Ala	aaa Lys	gga Gly 870	caa Gln	caa Gln	2766
gta Val	aaa Lys	tac Tyr 875	aca Thr	gtc Val	gag Glu	gaa Glu	tta Leu 880	Thr	aag Lys	gtc Val	aaa Lys	ggt Gly 885	tat Tyr	aca Thr	aca Thr	2814
cat His	gtg Val 890	Asp	aac Asn	aat Asn	gat Asp	atg Met 895	Gly	aac Asn	ttg Leu	att Ile	gtg Val 900	Thr	aat Asn	aaa Lys	tat Tyr	2862
acg Thr 905	Pro	gaa Glu	aca Thr	aca Thr	tca Ser 910	Ile	agt Ser	ggt Gly	gaa Glu	aaa Lys 915	Val	tgg Trp	gac Asp	gac Asp	aaa Lys 920	2910
gac Asp	aat Asn	caa Gln	gat Asp	ggt Gly 925	Lys	aga Arg	cca Pro	gaa Glu	aaa Lys 930	Val	agt Ser	gtg Val	aat Asn	tta Leu 935	ttg Leu	2958
gct Ala	aac Asn	gga Gly	gag Glu 940	Lys	gta Val	aaa Lys	acg Thr	tta Leu 945	Asp	gtg Val	aca Thr	tct Ser	gaa Glu 950	Thr	aac Asn	3006
tgg	aag	tac	gaa	ttt	aaa	gac	tta	ccg	aag	r tat	gat	gaa	gga	aag	aaa	3054

Trp	Lys	Tyr 955	Glu	Phe L	ys A		eu P 60	ro L	уѕ Ту	r As	p G1 96		у Lу	s Lys	
ata Ile	gaa Glu 970	tat Tyr	aca Thr	gtg a Val T	hr G	aa g lu A 175	at c sp H	ac g is V	ta aa al Ly	a ga s As 98	р Ту	c ac r Th	a ac r Th	a gac r Asp	3102
atc Ile 985	aac Asn.	ggt Gly	acg Thr	Thr I	ta a le T 90	icg a	ac a sn I	ag t ys T	at ac 'yr Th 99	r Pr	a go o Gl	ja ga .y Gl	g ac u Th	a tcg ir Ser 1000	3150
				aaa Lys 1005											3195
				gaa Glu 1020						tat Tyr	caa Gln	gat Asp	gga Gly	aaa Lys 1030	3240
				acg Thr 1035								aac Asn			3285
				gga Gly 1050					gca Ala 1055			caa Gln			3330
				gat Asp 1065								tat Tyr			3375
cat His	gtg Val	gat Asp	aac Asn	aat Asn 1080	gat Asp	atg Met	ggt Gly	aac Asn	ttg Leu 1085			aca Thr			3420
	-	_		aaa Lys 1095					atc Ile 1100			gaa Glu			3465
	_			cca Pro 1110					gat Asp 1115	_		aat Asn			3510
									aaa Lys 1130			aag Lys	_	-	3555
			_	aat Asn 1140					gaa Glu 1145			cta Leu			3600
				ggt Gly 1155					act Thr 1160			att Ile			3645
				ata Ile 1170					tta Leu 1175			_		aga Arg 1180	3690
ttt	aac	tca	taa	acca	ttat	aa t	tatt	ttta	t aga	taag	gct	attc	ttag	tt	3742

Phe Asn Ser

ctatgtataa tacatgtata ttaataggtc acttttaatc tgtatgtaag cagactaaga 3802 gtggcctttt aaacaaataa aaaaa 3827

Ile Ile Thr Pro Leu Phe Asn Lys Asn Glu Ala Phe Ala Ala Arg Asp 20 25 30

Ile Ser Ser Thr Asn Val Thr Asp Leu Thr Val Ser Pro Ser Lys Ile 35 40 45

Glu Asp Gly Gly Lys Thr Thr Val Lys Met Thr Phe Asp Asp Lys Asn 50 55

Gly Lys Ile Gln Asn Gly Asp Met Ile Lys Val Ala Trp Pro Thr Ser 65 70 75 80

Gly Thr Val Lys Ile Glu Gly Tyr Ser Lys Thr Val Pro Leu Thr Val 85 90 95

Lys Gly Glu Gln Val Gly Gln Ala Val Ile Thr Pro Asp Gly Ala Thr 100 $$ 105 $$ $$ $$ $$ $$

Ile Thr Phe Asn Asp Lys Val Glu Lys Leu Ser Asp Val Ser Gly Phe 115 125

Ala Glu Phe Glu Val Gln Gly Arg Asn Leu Thr Gln Thr Asn Thr Ser 130 135 140

Asp Asp Lys Val Ala Thr Ile Thr Ser Gly Asn Lys Ser Thr Asn Val 145 150 155 160

Thr Val His Lys Ser Glu Ala Gly Thr Ser Ser Val Phe Tyr Tyr Lys 165 170 175

Thr Gly Asp Met Leu Pro Glu Asp Thr Thr His Val Arg Trp Phe Leu 180 185 190

Asn Ile Asn Asn Glu Lys Ser Tyr Val Ser Lys Asp Ile Thr Ile Lys 195 200 205

Asp Gln Ile Gln Gly Gly Gln Gln Leu Asp Leu Ser Thr Leu Asn Ile 210 215 220

Asn Val Thr Gly Thr His Ser Asn Tyr Tyr Ser Gly Gln Ser Ala Ile 225 230 235 240

Thr Asp Phe Glu Lys Ala Phe Pro Gly Ser Lys Ile Thr Val Asp Asn 245 250 255

Thr Lys Asn Thr Ile Asp Val Thr Ile Pro Gln Gly Tyr Gly Ser Tyr 260 265 270

Asn Ser Phe Ser Ile Asn Tyr Lys Thr Lys Ile Thr Asn Glu Gln Gln 275 280 285

Lys Glu Phe Val Asn Asn Ser Gln Ala Trp Tyr Gln Glu His Gly Lys 290 295 300

Glu Glu Val Asn Gly Lys Ser Phe Asn His Thr Val His Asn Ile Asn 305 310 315 320

Ala Asn Ala Gly Ile Glu Gly Thr Val Lys Gly Glu Leu Lys Val Leu 325 330 335

Lys Gln Asp Lys Asp Thr Lys Ala Pro Ile Ala Asn Val Lys Phe Lys 340 345 350

Leu Ser Lys Lys Asp Gly Ser Val Val Lys Asp Asn Gln Lys Glu Ile 355 360 365

Glu Ile Ile Thr Asp Ala Asn Gly Ile Ala Asn Ile Lys Ala Leu Pro $370 \hspace{1cm} 375 \hspace{1cm} 380$

Ser Gly Asp Tyr Ile Leu Lys Glu Ile Glu Ala Pro Arg Pro Tyr Thr 385 390 395 400

Phe Asp Lys Asp Lys Glu Tyr Pro Phe Thr Met Lys Asp Thr Asp Asn 405 410 415

Gln Gly Tyr Phe Thr Thr Ile Glu Asn Ala Lys Ala Ile Glu Lys Thr 420 425 430

Lys Asp Val Ser Ala Gln Lys Val Trp Glu Gly Thr Gln Lys Val Lys 435 440 445

Pro Thr Ile Tyr Phe Lys Leu Tyr Lys Gln Asp Asp Asn Gln Asn Thr 450 455 460

Thr Pro Val Asp Lys Ala Glu Ile Lys Lys Leu Glu Asp Gly Thr Thr 465 470 475 480

Lys Val Thr Trp Ser Asn Leu Pro Glu Asn Asp Lys Asn Gly Lys Ala 485 490 495

Ile Lys Tyr Leu Val Lys Glu Val Asn Ala Gln Gly Glu Asp Thr Thr 500 505 510

Pro Glu Gly Tyr Thr Lys Lys Glu Asn Gly Leu Val Val Thr Asn Thr 515 520 525

Glu Lys Pro Ile Glu Thr Thr Ser Ile Ser Gly Glu Lys Val Trp Asp 530 535 540

Asp Lys Asp Asn Gln Asp Gly Lys Arg Pro Glu Lys Val Ser Val Asn 545 550 555 560

Leu Leu Ala Asn Gly Glu Lys Val Lys Thr Leu Asp Val Thr Ser Glu 565 570 575

Thr Asn Trp Lys Tyr Glu Phe Lys Asp Leu Pro Lys Tyr Asp Glu Gly 580 585 590

Lys Lys Ile Glu Tyr Thr Val Thr Glu Asp His Val Lys Asp Tyr Thr 595 600 605

Thr Asp Ile Asn Gly Thr Thr Ile Thr Asn Lys Tyr Thr Pro Gly Glu 610 620

Thr Ser Ala Thr Val Thr Lys Asn Trp Asp Asp Asn Asn Asn Gln Asp 625 630 635 640

Gly Lys Arg Pro Thr Glu Ile Lys Val Glu Leu Tyr Gln Asp Gly Lys 645 650 655

Ala Thr Gly Lys Thr Ala Ile Leu Asn Glu Ser Asn Asn Trp Thr His $660 \hspace{1.5cm} 665 \hspace{1.5cm} 670 \hspace{1.5cm}$

Thr Trp Thr Gly Leu Asp Glu Lys Ala Lys Gly Gln Gln Val Lys Tyr 675 680 685

Thr Val Glu Glu Leu Thr Lys Val Lys Gly Tyr Thr Thr His Val Asp 690 695 700

Asn Asn Asp Met Gly Asn Leu Ile Val Thr Asn Lys Tyr Thr Pro Glu 705 710 715 720

Thr Thr Ser Ile Ser Gly Glu Lys Val Trp Asp Asp Lys Asp Asn Gln 725 730 735

Asp Gly Lys Arg Pro Glu Lys Val Ser Val Asn Leu Leu Ala Asp Gly 740 745 750

Glu Lys Val Lys Thr Leu Asp Val Thr Ser Glu Thr Asn Trp Lys Tyr 755 760 765

Glu Phe Lys Asp Leu Pro Lys Tyr Asp Glu Gly Lys Lys Ile Glu Tyr 770 775 780

Thr Val Thr Glu Asp His Val Lys Asp Tyr Thr Thr Asp Ile Asn Gly 785 790 795 800

Thr Thr Ile Thr Asn Lys Tyr Thr Pro Gly Glu Thr Ser Ala Thr Val $805 \hspace{1.5cm} 810 \hspace{1.5cm} 815$

Thr Lys Asn Trp Asp Asp Asn Asn Asn Gln Asp Gly Lys Arg Pro Thr 820 825 830

Glu Ile Lys Val Glu Leu Tyr Gln Asp Gly Lys Ala Thr Gly Lys Thr 835 840 845

Ala Ile Leu Asn Glu Ser Asn Asn Trp Thr His Thr Trp Thr Gly Leu 850 855 860

Asp Glu Lys Ala Lys Gly Gln Gln Val Lys Tyr Thr Val Glu Glu Leu 865 870 875 880

Thr Lys Val Lys Gly Tyr Thr Thr His Val Asp Asn Asn Asp Met Gly 885 890 895

Asn Leu Ile Val Thr Asn Lys Tyr Thr Pro Glu Thr Thr Ser Ile Ser 900 905 910

Gly Glu Lys Val Trp Asp Asp Lys Asp Asn Gln Asp Gly Lys Arg Pro 915 920 925

- Glu Lys Val Ser Val Asn Leu Leu Ala Asn Gly Glu Lys Val Lys Thr 930 935 940
- Leu Asp Val Thr Ser Glu Thr Asn Trp Lys Tyr Glu Phe Lys Asp Leu 945 950 955 960
- Pro Lys Tyr Asp Glu Gly Lys Lys Ile Glu Tyr Thr Val Thr Glu Asp 965 970 975
- His Val Lys Asp Tyr Thr Thr Asp Ile Asn Gly Thr Thr Ile Thr Asn $980 \hspace{1.5cm} 985 \hspace{1.5cm} 990$
- Lys Tyr Thr Pro Gly Glu Thr Ser Ala Thr Val Thr Lys Asn Trp Asp 995 1000 1005
- Asp Asn Asn Asn Gln Asp Gly Lys Arg Pro Thr Glu Ile Lys Val 1010 1015 1020
- Glu Leu Tyr Gln Asp Gly Lys Ala Thr Gly Lys Thr Ala Ile Leu 1025 1030 1035
- Asn Glu Ser Asn Asn Trp Thr His Thr Trp Thr Gly Leu Asp Glu 1040 1045 1050
- Lys Ala Lys Gly Gln Gln Val Lys Tyr Thr Val Asp Glu Leu Thr $1055 \hspace{1.5cm} 1060 \hspace{1.5cm} 1065$
- Lys Val Asn Gly Tyr Thr Thr His Val Asp Asn Asn Asp Met Gly 1070 1075 1080
- Pro Ile Tyr Pro Glu Lys Pro Lys Asp Lys Thr Pro Pro Thr Lys 1100 1105
- Pro Asp His Ser Asn Lys Val Lys Pro Thr Pro Pro Asp Lys Pro 1115 1120 1125
- Ser Lys Val Asp Lys Asp Asp Gln Pro Lys Asp Asn Lys Thr Lys 1130 1135 1140
- Pro Glu Asn Pro Leu Lys Glu Leu Pro Lys Thr Gly Met Lys Ile 1145 1150 1155

Tyr Leu Ile Leu Arg Lys Arg Phe Asn Ser 1175 1180

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<220> <221> MISC_FEATURE <222> (6),.(7) <223> Xaa can be any amino acid

<400> 46

Ile Leu Val Xaa Tyr Xaa Xaa Leu Val

<210> 49 <211> 237 <212> PRT <213> Homo sapiens <400> 49

Met Thr Ser Glu Ile Thr Tyr Ala Glu Val Arg Phe Lys Asn Glu Phe $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Lys Ser Ser Gly Ile Asn Thr Ala Ser Ser Ala Ala Ser Lys Glu Arg

Thr Ala Pro His Lys Ser Asn Thr Gly Phe Pro Lys Leu Cys Ala 35 40 45

Ser Leu Leu Ile Phe Phe Leu Leu Ala Ile Ser Phe Phe Ile Ala 50 55 60

Phe Val Ile Phe Phe Gln Lys Tyr Ser Gln Leu Leu Glu Lys Lys Thr 65 70 75 80

Thr Lys Glu Leu Val His Thr Thr Leu Glu Cys Val Lys Lys Asn Met 85 90 95

Pro Val Glu Glu Thr Ala Trp Ser Cys Cys Pro Lys Asn Trp Lys Ser 100 105 110

Phe Ser Ser Asn Cys Tyr Phe Ile Ser Thr Glu Ser Ala Ser Trp Gln

115 120 125

Asp Ser Glu Lys Asp Cys Ala Arg Met Glu Ala His Leu Leu Val Ile 130 135 140

Asn Thr Gln Glu Glu Gln Asp Phe Ile Phe Gln Asn Leu Gln Glu Glu 145 150 155 160

Ser Ala Tyr Phe Val Gly Leu Ser Asp Pro Glu Gly Gln Arg His Trp 165 170 175

Gln Trp Val Asp Gln Thr Pro Tyr Asn Glu Ser Ser Thr Phe Trp His 180 185 190

Pro Arg Glu Pro Ser Asp Pro Asn Glu Arg Cys Val Val Leu Asn Phe
195 200 205

Arg Lys Ser Pro Lys Arg Trp Gly Trp Asn Asp Val Asn Cys Leu Gly 210 215 220

Pro Gln Arg Ser Val Cys Glu Met Met Lys Ile His Leu 225 230 235

<210> 53 <211> 32 <212> PRT <213> Homo sapiens <220> <221> MISC_FEATURE <222> (2)..(2) <223> Xaa is Ile or Leu

<220> <221> MISC_FEATURE <222> (3)..(3) <223> Xaa can be any amino
acid

<220> <221> MISC_FEATURE <222> (6)..(6) <223> Xaa can be any amino
acid

 $<\!220\!><\!221\!>$ MISC_FEATURE $<\!222\!>$ (8)..(25) $<\!223\!>$ Xaa can be any amino acid

<220> <221> MISC FEATURE <222> (26)..(26) <223> Xaa is Thr or Asn

<220> <221> MISC_FEATURE <222> (31)..(31) <223> Xaa is Ile or Val

<220> <221> MISC_FEATURE <222> (32)..(32) <223> Xaa is Lys or Arg

<400> 53

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu Tyr Ser Glu Xaa Xaa 20 25 30

<210> 54 < 211> 6 < 212> PRT < 213> Homo sapiens < 220> < 221>

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Xaa Glu Tyr Ser Glu Xaa 1 5

<210> 56 <211> 626 <212> PRT <213> Homo sapiens <400> 56

Met Ile Phe Leu Thr Ala Leu Pro Leu Phe Trp Ile Met Ile Ser Ala 1 5 10 15

Ser Arg Gly Gly His Trp Gly Ala Trp Met Pro Ser Ser Ile Ser Ala 20 25 30

Phe Glu Gly Thr Cys Val Ser Ile Pro Cys Arg Phe Asp Phe Pro Asp 35 40 45

Glu Leu Arg Pro Ala Val Val His Gly Val Trp Tyr Phe Asn Ser Pro 50 60

Tyr Pro Lys Asn Tyr Pro Pro Val Val Phe Lys Ser Arg Thr Gln Val 65 70 75 80

Val His Glu Ser Phe Gln Gly Arg Ser Arg Leu Leu Gly Asp Leu Gly 85 90 95

Leu Arg Asn Cys Thr Leu Leu Leu Ser Asn Val Ser Pro Glu Leu Gly 100 105 110

Gly Lys Tyr Tyr Phe Arg Gly Asp Leu Gly Gly Tyr Asn Gln Tyr Thr 115 120 125

Phe Ser Glu His Ser Val Leu Asp Ile Val Asn Thr Pro Asn Ile Val 130 135 140

Val Pro Pro Glu Val Val Ala Gly Thr Glu Val Glu Val Ser Cys Met 145 150 155 160

Val Pro Asp Asn Cys Pro Glu Leu Arg Pro Glu Leu Ser Trp Leu Gly 165 170 175

Glu Gly Thr Trp Val Gln Val Ser Leu Leu His Phe Val Pro Thr Arg 195 200 205

Glu Ala Asn Gly His Arg Leu Gly Cys Gln Ala Ser Phe Pro Asn Thr 210 215 220

Thr Leu Gln Phe Glu Gly Tyr Ala Ser Met Asp Val Lys Tyr Pro Pro 225 230 235 240

Val Ile Val Glu Met Asn Ser Ser Val Glu Ala Ile Glu Gly Ser His $245 \hspace{1.5cm} 250 \hspace{1.5cm} 255$

Val Ser Leu Leu Cys Gly Ala Asp Ser Asn Pro Pro Pro Leu Leu Thr 260 265 270

Trp Met Arg Asp Gly Thr Val Leu Arg Glu Ala Val Ala Glu Ser Leu 275 280 285

Leu Leu Glu Leu Glu Glu Val Thr Pro Ala Glu Asp Gly Val Tyr Ala 290 295 300

Cys Leu Ala Glu Asn Ala Tyr Gly Gln Asp Asn Arg Thr Val Gly Leu 305 310 315 320

Ser Val Met Tyr Ala Pro Trp Lys Pro Thr Val Asn Gly Thr Met Val 325 330 335

Ala Val Glu Gly Glu Thr Val Ser Ile Leu Cys Ser Thr Gln Ser Asn 340 345 350

Pro Asp Pro Ile Leu Thr Ile Phe Lys Glu Lys Gln Ile Leu Ser Thr 355 360 365

Val Ile Tyr Glu Ser Glu Leu Gln Leu Glu Leu Pro Ala Val Ser Pro 370 380

Glu Asp Asp Gly Glu Tyr Trp Cys Val Ala Glu Asn Gln Tyr Gly Gln 385 390 395 400

Arg Ala Thr Ala Phe Asn Leu Ser Val Glu Phe Ala Pro Val Leu Leu 405 410 415

Leu Glu Ser His Cys Ala Ala Ala Arg Asp Thr Val Gln Cys Leu Cys 420 425 430

Val Val Lys Ser Asn Pro Glu Pro Ser Val Ala Phe Glu Leu Pro Ser 435 440 445

Arg Asn Val Thr Val Asn Glu Ser Glu Arg Glu Phe Val Tyr Ser Glu 450 455 460

Arg Ser Gly Leu Val Leu Thr Ser Ile Leu Thr Leu Arg Gly Gln Ala 465 470 475 480

Gln Ala Pro Pro Arg Val Ile Cys Thr Ala Arg Asn Leu Tyr Gly Ala 485 490 495

Lys Ser Leu Glu Leu Pro Phe Gln Gly Ala His Arg Leu Met Trp Ala 500 505 510

Lys Ile Gly Pro Val Gly Ala Val Val Ala Phe Ala Ile Leu Ile Ala 515 520 525

Ile Val Cys Tyr Ile Thr Gln Thr Arg Arg Lys Lys Asn Val Thr Glu 530 535 540

Ser Pro Ser Phe Ser Ala Gly Asp Asn Pro Pro Val Leu Phe Ser Ser 545 550 555 560

Asp Phe Arg Ile Ser Gly Ala Pro Glu Lys Tyr Glu Ser Glu Arg Arg 565 570 575

Leu Gly Ser Glu Arg Arg Leu Leu Gly Leu Arg Gly Glu Pro Pro Glu 580 585 590

Leu Asp Leu Ser Tyr Ser His Ser Asp Leu Gly Lys Arg Pro Thr Lys 595 600

Asp Ser Tyr Thr Leu Thr Glu Glu Leu Ala Glu Tyr Ala Glu Ile Arg 610 620

Val Lys 625

Gln Ile Glu Thr Gly Pro Phe Leu Glu Ala Val Ser His Leu Pro Pro 20 25 30

Phe Phe Asp Cys Leu Gly Ser Pro Val Phe Thr Pro Ile Lys Ala Asp 35 40 45

Ile Ser Gly Asn Ile Thr Lys Ile Lys Ala Val Tyr Asp Thr Asn Pro 50 55 60

Ala Lys Phe Arg Thr Leu Gln Asn Ile Leu Glu Val Glu Lys Glu Met 65 70 75 80

Tyr Gly Ala Glu Trp Pro Lys Val Gly Ala Thr Leu Ala Leu Met Trp 85 90 95

Leu Lys Arg Gly Leu Arg Phe Ile Gln Val Phe Leu Gln Ser Ile Cys 100 105 110

Asp Gly Glu Arg Asp Glu Asn His Pro Asn Leu Ile Arg Val Asn Ala 115 120 125

Thr Lys Ala Tyr Glu Met Ala Leu Lys Lys Tyr His Gly Trp Ile Val 130 135 140

Gln Lys Ile Phe Gln Ala Ala Leu Tyr Ala Ala Pro Tyr Lys Ser Asp 145 150 155 160

Phe Leu Lys Ala Leu Ser Lys Gly Gln Asn Val Thr Glu Glu Glu Cys 165 170 175

Leu Glu Lys Ile Arg Leu Phe Leu Val Asn Tyr Thr Ala Thr Ile Asp 180 185 190

Val Ile Tyr Glu Met Tyr Thr Gln Met Asn Ala Glu Leu Asn Tyr Lys 195 200 205

Val

<210> 64 <211> 276 <212> PRT <213> Homo sapiens <400> 64

Met Gly Asn Ser Met Lys Ser Thr Pro Ala Pro Ala Glu Arg Pro Leu $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Pro Asn Pro Glu Gly Leu Asp Ser Asp Phe Leu Ala Val Leu Ser Asp 20 25 30

Tyr Pro Ser Pro Asp Ile Ser Pro Pro Ile Phe Arg Arg Gly Glu Lys
35 40 45

Leu Arg Val Ile Ser Asp Glu Gly Gly Trp Trp Lys Ala Ile Ser Leu 50 55 60

Ser Thr Gly Arg Glu Ser Tyr Ile Pro Gly Ile Cys Val Ala Arg Val 65 70 75 80

Tyr His Gly Trp Leu Phe Glu Gly Leu Gly Arg Asp Lys Ala Glu Glu 85 90 95

Leu Leu Gln Leu Pro Asp Thr Lys Val Gly Ser Phe Met Ile Arg Glu 100 105 110

Ser Glu Thr Lys Lys Gly Phe Tyr Ser Leu Ser Val Arg His Arg Gln $1\dot{1}5$ 120 125

Val Lys His Tyr Arg Ile Phe Arg Leu Pro Asn Asn Trp Tyr Tyr Ile 130 135 140

Ser Pro Arg Leu Thr Phe Gln Cys Leu Glu Asp Leu Val Asn His Tyr 145 150 155 160

Ser Glu Val Ala Asp Gly Leu Cys Cys Val Leu Thr Thr Pro Cys Leu 165 170 175

Thr Gln Ser Thr Ala Ala Pro Ala Val Arg Ala Ser Ser Ser Pro Val 180 185 190

Thr Leu Arg Gln Lys Thr Val Asp Trp Arg Arg Val Ser Arg Leu Gln 195 200 205

Glu Asp Pro Glu Gly Thr Glu Asn Pro Leu Gly Val Asp Glu Ser Leu 210 $$ 220

Phe Ser Tyr Gly Leu Arg Glu Ser Ile Ala Ser Tyr Leu Ser Leu Thr 225 230 235 240

Ser Glu Asp Asn Thr Ser Phe Asp Arg Lys Lys Ser Ile Ser Leu 245 250 255

Met Tyr Gly Gly Ser Lys Arg Lys Ser Ser Phe Phe Ser Ser Pro Pro 260 265 270

Tyr Phe Glu Asp 275

Ser Cys

Ser Cys

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Phe Ile Gln Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Val Leu

67 <211> 1784 <212> DNA <213> Staphylococcus aureus <220>

	65					70					75					
						aat Asn										468
						aaa Lys										516
						aac Asn										564
				_		agc Ser		_						-	_	612
						caa Gln 150		_		_	_					660
					_	ttc Phe		_		_		_				708
						ggt Gly										756
						ttg Leu										804
	_	_	Lys		-	aac Asn					-				-	852
						tta Leu 230										900
						aaa Lys										948
tta Leu	gca Ala	gaa Glu	gct Ala	aaa Lys 260	aag Lys	cta Leu	aat Asn	gat Asp	gct Ala 265	caa Gln	gca Ala	cca Pro	aaa Lys	gct Ala 270	gac Asp	996
					-	caa Gln			_			_				1044
						gaa Glu										1092
						agc Ser										1140

	305					310					315					
		_	_		-				_	gac Asp 330			-			1188
										gac Asp						1236
	_	_			_				_	gaç Asp			_			1284
	_	-			_				_	gac Asp			_		J J	1332
	-	_							-	gat Asp			_			1380
	_	_							_	gac Asp 410			_		~ ~	1428
										gat Asp						1476
										att Ile						1524
		_	-			-	_	_		aaa Lys		_				1572
_						_		_	_	gat Asp	_	_				1620
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ttca	atcgo	gta d	caact	gtat	it to	ggtg	gatta	a tca	atta	gcct	tag	gtgca	agc (gtta	ttagct	1735
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- Ala Ala As
n Ala Gl
n His Asp Glu Ala Gl
n Gln As
n Ala Phe Tyr 35 4045
- Gln Val Leu Asn Met Pro Asn Leu Asn Ala Asp Gln Arg Asn Gly Phe 50 60
- Ile Gln Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Val Leu Gly 65 70 75 80
- Glu Ala Gln Lys Leu Asn Asp Ser Gln Ala Pro Lys Ala Asp Ala Gln 85 90 95
- Gln Asn Asn Phe Asp Lys Asp Gln Gln Ser Ala Phe Tyr Glu Ile Leu 100 105 110
- Asn Met Pro Asn Leu Asn Glu Ala Gln Arg Asn Gly Phe Ile Gln Ser 115 120 125
- Leu Lys Asp Asp Pro Ser Gln Ser Thr Asn Val Leu Gly Glu Ala Lys 130 140
- Lys Leu Asn Glu Ser Gln Ala Pro Lys Ala Asp Asn Asn Phe Asn Lys 145 150 155 160
- Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu Asn Met Pro Asn Leu Asn 165 170 175
- Glu Glu Gln Arg Asn Gly Phe Ile Gln Ser Leu Lys Asp Asp Pro Ser 180 185 190
- Gln Ser Ala Asn Leu Leu Ser Glu Ala Lys Lys Leu Asn Glu Ser Gln 195 200 205
- Ala Pro Lys Ala Asp Asn Lys Phe Asn Lys Glu Gln Gln Asn Ala Phe 210 215 220
- Tyr Glu Ile Leu His Leu Pro Asn Leu Asn Glu Glu Gln Arg Asn Gly 225 230 235 240
- Phe Ile Gln Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu Leu 245 250 255
- Ala Glu Ala Lys Lys Leu As
n Asp Ala Gl
n Ala Pro Lys Ala Asp As
n 260 265 270

Lys Phe Asn Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu His Leu 275 280 285

Pro Asn Leu Thr Glu Glu Gln Arg Asn Gly Phe Ile Gln Ser Leu Lys 290 295 300

Asp Asp Pro Ser Val Ser Lys Glu Ile Leu Ala Glu Ala Lys Lys Leu 305 310 315 320

Asn Asp Ala Gln Ala Pro Lys Glu Glu Asp Asn Asn Lys Pro Gly Lys 325 330 335

Glu Asp Asn Asn Lys Pro Gly Lys Glu Asp Asn Asn Lys Pro Gly Lys 340 345 350

Glu Asp Asn Asn Lys Pro Gly Lys Glu Asp Asn Asn Lys Pro Gly Lys 355 360 365

Glu Asp Asn Asn Lys Pro Gly Lys Glu Asp Gly Asn Lys Pro Gly Lys 370 375 380

Glu Asp Asn Lys Lys Pro Gly Lys Glu Asp Gly Asn Lys Pro Gly Lys 385 390 395 400

Glu Asp Asn Lys Lys Pro Gly Lys Glu Asp Gly Asn Lys Pro Gly Lys
405
410
415

Glu Asp Gly Asn Lys Pro Gly Lys Glu Asp Gly Asn Gly Val His Val 420 425 430

Val Lys Leu Gly Asp Thr Val Asn Asp Ile Ala Lys Ala Asn Gly Thr 435 440 445

Thr Ala Asp Lys Ile Ala Ala Asp Asn Lys Leu Ala Asp Lys Asn Met 450 455 460

Ile Lys Leu Gly Gln Glu Leu Val Val Asp Lys Lys Gln Pro Gln Thr 465 470 475 480

Met Gln Ser

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											tac Tyr					279
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Arg	Lon	T 0	_													
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Trp			20					25			Tyr		30			
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Gly Leu 65	Val Asp 50 Asp	Lys 35 Ile Val	Trp Leu Val	Met Pro	Arg Asn Gly	Gly Ala 55 Glu	Glu 40 Asp	Gln Glu Ala	Glu Thr	Gln Trp Leu 75	Gln Tyr 60	Gly 45 Leu Cys	30 Thr Arg	Gln Ala Val	Pro Thr Lys	
Gly Leu 65	Val Asp 50 Asp Ser	Lys 35 Ile Val Ser	20 Trp Leu Val Leu	Met Pro Ala Glu 85	Arg Asn Gly	Gly Ala 55 Glu Gln 212>	Glu 40 Asp Ala Asp	Glu Ala Ile	Glu Thr Gly Val	Gln Trp Leu 75	Gln Tyr 60 Ser	Gly 45 Leu Cys	30 Thr Arg	Gln Ala Val	Pro Thr Lys 80	CDS
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gaacaageee gtggaacetg gaageeagag aacaceateg etgtaaagga aagaaactga	120
agettttete ggageetate tgggetgete aaacttteag aatetaeega ee atg Met 1	175
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tggcaggagt ctccctgtcc ccctactttc tcctcag gag cgt gtg ctt ggc ttg Glu Arg Val Leu Gly Leu 5	290
ttg ctg ttg ctt ctg gtg cac gcc tct ccc gcc cca cca gag ccc tgc Leu Leu Leu Leu Val His Ala Ser Pro Ala Pro Pro Glu Pro Cys 10 15 20	338
gag cta gac gag gaa agt tgc tcc tgc aac ttc tca gat ccg aag cca Glu Leu Asp Glu Glu Ser Cys Ser Cys Asn Phe Ser Asp Pro Lys Pro 25 30 35	386
gat tgg tcc agc gct ttc aat tgt ttg ggg gcg gca gat gtg gaa ttg Asp Trp Ser Ser Ala Phe Asn Cys Leu Gly Ala Ala Asp Val Glu Leu 40 45 50 55	434
tac ggc ggc cgc agc ctg gaa tac ctt cta aag cgt gtg gac acg Tyr Gly Gly Gly Arg Ser Leu Glu Tyr Leu Leu Lys Arg Val Asp Thr 60 65 70	482
gaa gca gat ctg ggg cag ttc act gat att atc aag tct ctg tcc tta Glu Ala Asp Leu Gly Gln Phe Thr Asp Ile Ile Lys Ser Leu Ser Leu 75 80 85	530
aag cgg ctt acg gtg cgg gcc gcg cgg att cct agt cgg att cta ttc Lys Arg Leu Thr Val Arg Ala Ala Arg Ile Pro Ser Arg Ile Leu Phe 90 95 100	578
gga gcc ctg cgt gtg ctc ggg att tcc ggc ctc cag gaa ctg act ctt Gly Ala Leu Arg Val Leu Gly Ile Ser Gly Leu Gln Glu Leu Thr Leu 105 110 115	626
gaa aat ctc gag gta acc ggc acc gcg ccg cca ccg ctt ctg gaa gcc Glu Asn Leu Glu Val Thr Gly Thr Ala Pro Pro Pro Leu Leu Glu Ala 120 125 130 135	674
acc gga ccc gat ctc aac atc ttg aac ctc cgc aac gtg tcg tgg gca Thr Gly Pro Asp Leu Asn Ile Leu Asn Leu Arg Asn Val Ser Trp Ala 140 145 150	722
aca agg gat gcc tgg ctc gca gaa ctg cag cag tgg cta aag cct gga Thr Arg Asp Ala Trp Leu Ala Glu Leu Gln Gln Trp Leu Lys Pro Gly 155 160 165	770
ctc aag gta ctg agt att gcc caa gca cac tca ctc aac ttt tcc tgc Leu Lys Val Leu Ser Ile Ala Gln Ala His Ser Leu Asn Phe Ser Cys 170 175 180	818
gaa cag gtc cgc gtc ttc cct gcc ctc tcc acc tta gac ctg tct gac. Glu Gln Val Arg Val Phe Pro Ala Leu Ser Thr Leu Asp Leu Ser Asp	866

185	190	195	
aat cct gaa ttg ggc gag Asn Pro Glu Leu Gly Glu 200 205			914
aag ttc ccg acc ctc caa Lys Phe Pro Thr Leu Gln 220			962
acg ccc agc ggc gtg tgc Thr Pro Ser Gly Val Cys 235			1010
caa gga cta gac ctt agt Gln Gly Leu Asp Leu Ser 250		3 3 33 3	1058
ccg agt tgt gac tgg ccc Pro Ser Cys Asp Trp Pro 265	2 2	3	1106
act ggg ctg aag cag gta Thr Gly Leu Lys Gln Val 280 285			1154
ctg gat ctc agt tac aac Leu Asp Leu Ser Tyr Asn 300			1202
ctg ccc caa gtg ggg aac Leu Pro Gln Val Gly Asn 315	-	2 2	1250
tct gaa tcc cac tcg gag Ser Glu Ser His Ser Glu 330	2		1298
gct cca tca tcc caa gca Ala Pro Ser Ser Gln Ala 345			1346
cta gga gat cgc ctc ttt Leu Gly Asp Arg Leu Phe 360 365		atcctc ctggtttctg	1397
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ttaaccctca ggatccacac g	gaaggggaa aactcactcc	tgaaagttgt ccatctgtgc	1577
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tggcctcaaa ctcataaaga t	caagategg eetgeeteta	cctccaaatg ctctggttaa	1757
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1873

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Asn Phe Ser Asp Pro Lys Pro Asp Trp Ser Ser Ala Phe Asn Cys Leu 35 40 45

Gly Ala Ala Asp Val Glu Leu Tyr Gly Gly Gly Arg Ser Leu Glu Tyr 50 55 60

Leu Leu Lys Arg Val Asp Thr Glu Ala Asp Leu Gly Gln Phe Thr Asp 65 70 75 80

Ile Ile Lys Ser Leu Ser Leu Lys Arg Leu Thr Val Arg Ala Arg 85 90 95

Ile Pro Ser Arg Ile Leu Phe Gly Ala Leu Arg Val Leu Gly Ile Ser 100 105 110

Gly Leu Gl
n Glu Leu Thr Leu Glu Asn Leu Glu Val Thr Gly Thr Ala
 115 120 125

Leu Arg Asn Val Ser Trp Ala Thr Arg Asp Ala Trp Leu Ala Glu Leu 145 150 155 160

Gln Gln Trp Leu Lys Pro Gly Leu Lys Val Leu Ser Ile Ala Gln Ala 165 170 175

His Ser Leu Asn Phe Ser Cys Glu Gln Val Arg Val Phe Pro Ala Leu 180 185 190

Ser Thr Leu Asp Leu Ser Asp Asn Pro Glu Leu Gly Glu Arg Gly Leu 195 200 205

Ile Ser Ala Leu Cys Pro Leu Lys Phe Pro Thr Leu Gln Val Leu Ala 210 215 220

Leu Arg Asn Ala Gly Met Glu Thr Pro Ser Gly Val Cys Ser Ala Leu 225 230 235 240

Ala Ala Ala Arg Val Gln Leu Gln Gly Leu Asp Leu Ser His Asn Ser 245 250 255

Leu Arg Asp Ala Ala Gly Ala Pro Ser Cys Asp Trp Pro Ser Gln Leu 260 265 270

Asn Ser Leu Asn Leu Ser Phe Thr Gly Leu Lys Gln Val Pro Lys Gly 275 280 285

Leu Pro Ala Lys Leu Ser Val Leu Asp Leu Ser Tyr Asn Arg Leu Asp 290 295 300

Lys Gly Asn Pro Phe Leu Asp Ser Glu Ser His Ser Glu Lys Phe Asn 325 330 335

Ser Gly Val Val Thr Ala Gly Ala Pro Ser Ser Gln Ala Val Ala Leu 340 345 350

Ser Gly Thr Leu Ala Leu Leu Gly Asp Arg Leu Phe Val 355 360 365

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Ser Arg Thr Glu Ala Ala Asp Leu Cys Gln Ala Phe Asn Ser Thr Leu 50 55 60

Pro Thr Met Asp Gln Met Lys Leu Ala Leu Ser Lys Gly Phe Glu Thr 65 70 75 80

Cys Arg Tyr Gly Phe Ile Glu Gly Asn Val Val Ile Pro Arg Ile His

85 90 95

Pro Asn Ala Ile Cys Ala Ala Asn His Thr Gly Val Tyr Ile Leu Val 100 105 110

Thr Ser Asn Thr Ser His Tyr Asp Thr Tyr Cys Phe Asn Ala Ser Ala 115 120 125

Asp Gly Pro Val Thr Ile Thr Ile Val Asn Arg Asp Gly Thr Arg Tyr 145 150 155 160

Ser Lys Lys Gly Glu Tyr Arg Thr His Gln Glu Asp Ile Asp Ala Ser 165 170 175

Asn Ile Ile Asp Asp Asp Val Ser Ser Gly Ser Thr Ile Glu Lys Ser 180 185 190

Thr Pro Glu Ser Tyr Ile Leu His Thr Tyr Leu Pro Thr Glu Gln Pro 195 200 205

Thr Gly Asp Gln Asp Asp Ser Phe Phe Ile Arg Ser Thr Leu Ala Thr 210 215 220

Arg Asp Arg Asp Ser Ser Lys Asp Ser Arg Gly Ser Ser Arg Thr Val 225 230 235 240

Thr His Gly Ser Glu Leu Ala Gly His Ser Ser Ala Asn Gln Asp Ser 245 250 255

Gly Val Thr Thr Ser Gly Pro Met Arg Arg Pro Gln Ile Pro Glu 260 265 270

Trp Leu Ile Ile Leu Ala Ser Leu Leu Ala Leu Ala Leu Ile Leu Ala 275 280 285

Val Cys Ile Ala Val Asn Ser Arg Arg Cys Gly Gln Lys Lys 290 295 300

Leu Val Ile Asn Gly Gly Asn Gly Thr Val Glu Asp Arg Lys Pro Ser 305 310 315 320

Glu Leu Asn Gly Glu Ala Ser Lys Ser Gln Glu Met Val His Leu Val

325 330 335

Asn Lys Glu Pro Ser Glu Thr Pro Asp Gln Cys Met Thr Ala Asp Glu 340 345 350

Thr Arg Asn Leu Gln Ser Val Asp Met Lys Ile Gly Val 355 360 365

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Val Val Val Phe Trp Glu Tyr Val Asn Ser Pro Asp Gly Ser Phe 20 25 30

Leu Trp Ile Tyr His Thr Lys Ile Pro Glu Val Gly Glu Asn Arg Trp 35 40 45

Gln Lys Asp Trp Trp Phe Pro Ser Trp Phe Lys Asn Gly Thr His Ser 50 55 60

Tyr Gln Glu Asp Asn Val Glu Gly Arg Arg Glu Lys Gly Arg Asn Gly 65 70 75 80

Asp Arg Ile Glu Glu Pro Gln Leu Trp Asp Trp Phe Asn Pro Lys Asn 85 90 95

Arg Pro Asp Val Leu Thr Val Thr Pro Trp Lys Ala Pro Ile Val Trp 100 105 110

Glu Gly Thr Tyr Asp Thr Ala Leu Leu Glu Lys Tyr Tyr Ala Thr Gln
115 120 125

Lys Leu Thr Val Gly Leu Thr Val Phe Ala Val Gly Lys Tyr Ile Glu 130 135 140

His Tyr Leu Glu Asp Phe Leu Glu Ser Ala Asp Met Tyr Phe Met Val 145 150 155 160

Gly His Arg Val Ile Phe Tyr Val Met Ile Asp Asp Thr Ser Arg Met 165 . 170 175

Pro Val Val His Leu Asn Pro Leu His Ser Leu Gln Val Phe Glu Ile 180 185 190

Arg	Ser	Glu 195	Lys	Arg	'l'rp	GIN	200	116	ser	мец	мес	205	Mec	цуз	1111		
	Gly 210	Glu	His	Ile	Leu	Ala 215	His	Ile	Gln	His	Glu 220	Val	Asp	Phe	Leu		
Phe 225	Cys	Met	Asp	Val	Asp 230	Gln	Val	Phe	Gln	Asp 235	Asn	Phe	Gly	Val	Glu 240		
Thr	Leu	Gly	Gln	Leu 245	Val	Ala	Gln	Leu	Gln 250	Ala	Trp	Trp	Tyr	Lys 255	Ala		
Ser	Pro	Glu	Lys 260	Phe	Thr	Tyr	Glu	Arg 265	Arg	Glu	Leu	Ser	Ala 270	Ala	Tyr		
Ile	Pro	Phe 275	Gly	Glu	Gly	Asp	Phe 280	Tyr	Tyr	His	Ala	Ala 285	Ile	Phe	Gly		
Gly	Thr 290		Thr	His	Ile	Leu 295	Asn	Leu	Thr	Arg	Glu 300	Cys	Phe	Lys	Gly		
Ile 305	Leu	Gln	Asp	Lys	Lys 310	His	Asp	Ile	Glu	Ala 315	Gln	Trp	His	Asp	Glu 320		
Ser	His	Leu	Ąsn	Lys 325		Phe	Leu	Phe	Asn 330	Lys	s Pro	Thr	Lys	335	Leu		
Ser	Pro	Glu	Tyr 340		: Trp	Asp	туг	Gln 345	Ile	e Gly	/ Leu	Pro	Ser 350	Asp	lle		
Lys	Sei	Val 355		s Val	L Alá	Trp	Glr 360	n Thr	Lys	s Glı	1 Туі	Asr 365	ı Let	ı Val	. Arg		
Asn	Ası 370	n Val	L														
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act Thi	t cg r Ar	a cto g Le	c tc u Se	c at r Il	c ct e Le	a ga u Gl	g ga u As	c cto p Le	g aa u As	n							7

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	150					155					160					
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aca Thr	aca Thr	aat Asn	gcc Ala	gtt Val 185	aat Asn	att Ile	gcg Ala	atg Met	cgt Arg 190	cac His	gca Ala	acc Thr	acc Thr	ccc Pro 195	aat Asn	693
ttt Phe	tca Ser	tcg Ser	gcg Ala 200	ctt Leu	aat Asn	att Ile	act Thr	agc Ser 205	ggc Gly	aat Asn	gaa Glu	aat Asn	ggt Gly 210	agt Ser	gca Ala	741
atg Met	cag Gln	cta Leu 215	cga Arg	ggg Gly	tca Ser	gaa Glu	aaa Lys 220	gcg Ala	cta Leu	gga Gly	acg Thr	cta Leu 225	aaa Lys	att Ile	act Thr	789
cat His	gag Glu 230	aac Asn	cca Pro	agt Sér	att Ile	gga Gly 235	gcg Ala	gat Asp	tat Tyr	gat Asp	aaa Lys 240	aat Asn	gcg Ala	gca Ala	cgt Arg	837
tat Tyr 245	cca Pro	ttg Leu	ata Ile	ttg Leu	tca Ser 250	aaa Lys	aga Arg	caa Gln	aac Asn	ggt Gly 255	gca Ala	gga Gly	aca Thr	gcc Ala	gct Ala 260	885
cag Gln	gga Gly	atc Ile	Tyr	att Ile 265	Asn	tca Ser	acc Thr	tca Ser	ggc Gly 270	acg Thr	aca Thr	ggg Gly	aag Lys	ttg Leu 275	ctt Leu	933
agg Arg	att Ile	aga Arg	aac Asn 280	Leu	agt Ser	gat Asp	gat Asp	aag Lys 285	Phe	tac Tyr	gtc Val	aag Lys	tct Ser 290	Asp	ggt Gly	981
ggt Gly	ttt Phe	tat Tyr 295	Ala	aag Lys	gaa Glu	act Thr	tcg Ser 300	Gln	att Ile	gat Asp	ggç Gly	aac Asn 305	Leu	aaa Lys	ctc Leu	1029
aag Lys	gac Asp 310	Pro	aca Thr	gcg Ala	aat Asn	gat Asp 315	His	gcg Ala	gca Ala	acc Thr	aaa Lys 320	Ala	tat Tyr	gta Val	gat Asp	1077
aaa Lys 325	Āla	att Ile	tct Ser	gaç Glu	tta Leu 330	Lys	aaa Lys	cto Leu	ata Ile	cta Lev 335	ı Lys	aaa Lys	tag	I		1119
att	aagg	jagg	ataa	atga	igc a	ıgaga	ccca	aa ca	ictta	tttt	aga	cgaç	jtca	aacc	ctcgtt	a 1179
ttç	ggtaa	ıgga	tgga	cgto	J											119
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Ala	a Glı	ı Tr <u>p</u>	Ala	a Ar	g Sei	r Asp	o Va	l Ile	e Lei	ı Leı	ı Glı	ı Se	r Gli 30	u Ile	e Gly	

Phe Glu Thr Asp Thr Gly Phe Ala Arg Ala Gly Asp Gly His Asn Arg 35 40 45

Phe Ser Asp Leu Gly Tyr Ile Ser Pro Leu Asp Tyr Asn Leu Leu Thr 50 55 60

Asn Lys Pro Asn Ile Asp Gly Leu Ala Thr Lys Val Glu Thr Ala Gln 65 70 75 80

Lys Leu Gln Gln Lys Ala Asp Lys Glu Thr Val Tyr Thr Lys Ala Glu 85 90 95

Ser Lys Gln Glu Leu Asp Lys Lys Leu Asn Leu Lys Gly Gly Val Met
100 105 110

Thr Gly Gln Leu Lys Phe Lys Pro Ala Ala Thr Val Ala Tyr Ser Ser 115 120 125

Ser Thr Gly Gly Ala Val Asn Ile Asp Leu Ser Ser Thr Arg Gly Ala 130 135 140

Gly Val Val Val Tyr Ser Asp Asn Asp Thr Ser Asp Gly Pro Leu Met 145 150 155 160

Ser Leu Arg Thr Gly Lys Glu Thr Phe Asn Gln Ser Ala Leu Phe Val 165 170 175

Asp Tyr Lys Gly Thr Thr Asn Ala Val Asn Ile Ala Met Arg His Ala 180 185 190

Thr Thr Pro Asn Phe Ser Ser Ala Leu Asn Ile Thr Ser Gly Asn Glu

Asn Gly Ser Ala Met Gln Leu Arg Gly Ser Glu Lys Ala Leu Gly Thr 210 215 220

Leu Lys Ile Thr His Glu Asn Pro Ser Ile Gly Ala Asp Tyr Asp Lys 225 230 235 240

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Gly Thr Ala Ala Gln Gly Ile Tyr Ile Asn Ser Thr Ser Gly Thr Thr $260 \hspace{1.5cm} 265 \hspace{1.5cm} 270 \hspace{1.5cm}$

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n His Asp Glu Ala Gl
n Gln As
n Ala Phe Tyr 35 40 45
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- Val Lys Pro Gly Asp Thr Val Asn Asp Ile Ala Lys Ala Asn Gly Thr 420 425 430
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	_			-	_			***	_		**	_		gaa Glu		250

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Val Gln Leu Ala His His Phe Ser Glu Pro Glu Ile Thr Leu Ile Ile

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85

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5185

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- Thr Asp Trp Val Ala Ile Thr Leu Tyr Ala Cys Asp Ser Lys Ser Glu 85 90 95
- Phe Gln Lys Trp Glu Cys Lys Asn Asp Thr Leu Leu Gly Ile Lys Gly 100 105 110
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- Leu Tyr Lys Gly Ser Gly Leu Trp Ser Arg Trp Lys Ile Tyr Gly Thr 130 135 140
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- Lys Trp Tyr Ala Asp Cys Thr Ser Ala Gly Arg Ser Asp Gly Trp Leu 180 185 190
- Trp Cys Gly Thr Thr Thr Asp Tyr Asp Thr Asp Lys Leu Phe Gly Tyr 195 200 205
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Ile Thr Glu Ile His Glu Gln Thr Tyr Leu Thr Gly Leu Thr Ser Ser 260 265 270

Leu Thr Ser Gly Leu Trp Ile Gly Leu Asn Ser Leu Ser Phe Asn Ser 275 280 285

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Pro Leu Gly Tyr Ile Cys Lys Met Lys Ser Arg Ser Gln Gly Pro Glu 485 490 495

Ile Val Glu Val Glu Lys Gly Cys Arg Lys Gly Trp Lys Lys His His 500 505 510

Phe Tyr Cys Tyr Met Ile Gly His Thr Leu Ser Thr Phe Ala Glu Ala 515 520 525

Asn Gln Thr Cys Asn Asn Glu Asn Ala Tyr Leu Thr Thr Ile Glu Asp 530 535 540

Arg Tyr Glu Gln Ala Phe Leu Thr Ser Phe Val Gly Leu Arg Pro Glu 545 550 555 560

Lys Tyr Phe Trp Thr Gly Leu Ser Asp Ile Gln Thr Lys Gly Thr Phe 565 570 575

Gln Trp Thr Ile Glu Glu Glu Val Arg Phe Thr His Trp Asn Ser Asp 580 585 590

Met Pro Gly Arg Lys Pro Gly Cys Val Ala Met Arg Thr Gly Ile Ala $595 \hspace{1.5cm} 600 \hspace{1.5cm} 605$

Gly Gly Leu Trp Asp Val Leu Lys Cys Asp Glu Lys Ala Lys Phe Val 610 615 620

Cys Lys His Trp Ala Glu Gly Val Thr His Pro Pro Lys Pro Thr Thr 625 630 635 640

Thr Pro Glu Pro Lys Cys Pro Glu Asp Trp Gly Ala Ser Ser Arg Thr 645 650 655

Ser Leu Cys Phe Lys Leu Tyr Ala Lys Gly Lys His Glu Lys Lys Thr 660 665 670

Trp Phe Glu Ser Arg Asp Phe Cys Arg Ala Leu Gly Gly Asp Leu Ala 675 680 685

Ser Ile Asn Asn Lys Glu Glu Gln Gln Thr Ile Trp Arg Leu Ile Thr 690 695 700

Ala Ser Gly Ser Tyr His Lys Leu Phe Trp Leu Gly Leu Thr Tyr Gly 705 710 715 720

Ser Pro Ser Glu Gly Phe Thr Trp Ser Asp Gly Ser Pro Val Ser Tyr 725 730 735

Glu Asn Trp Ala Tyr Gly Glu Pro Asn Asn Tyr Gln Asn Val Glu Tyr
740 745 750

Cys Gly Glu Leu Lys Gly Asp Pro Thr Met Ser Trp Asn Asp Ile Asn 755 760 765

Cys Glu His Leu Asn Asn Trp Ile Cys Gln Ile Gln Lys Gly Gln Thr 770 775 780

Pro Lys Pro Glu Pro Thr Pro Ala Pro Gln Asp Asn Pro Pro Val Thr 785 790 795 800

Glu Asp Gly Trp Val Ile Tyr Lys Asp Tyr Gln Tyr Tyr Phe Ser Lys 805 810 815

Glu Lys Glu Thr Met Asp Asn Ala Arg Ala Phe Cys Lys Arg Asn Phe 820 825 830

Lys Tyr Val Asn Arg Asn Asp Ala Gln Ser Ala Tyr Phe Ile Gly Leu 850 860

Leu Ile Ser Leu Asp Lys Lys Phe Ala Trp Met Asp Gly Ser Lys Val 865 870 875 880

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Glu Asn Cys Val Thr Met Tyr Ser Asn Ser Gly Phe Trp Asn Asp Ile $900 \hspace{1.5cm} 905 \hspace{1.5cm} 910$

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- Cys Lys Glu Gly Trp Asn Phe Tyr Ser Asn Lys Cys Phe Lys Ile Phe 945 950 955 960
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- Cys Ile Gly Phe Gly Gly Asn Leu Val Ser Ile Gln Asn Glu Lys Glu 980 985 990
- Gln Ala Phe Leu Thr Tyr His Met Lys Asp Ser Thr Phe Ser Ala Trp 995 1000 1005
- Thr Gly Leu Asn Asp Val Asn Ser Glu His Thr Phe Leu Trp Thr 1010 1015 1020
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- Val Ile Ile Gly Gly Ala Ser Asn Glu Ala Gly Lys Trp Met Asp 1055 1060 1065
- Asp Thr Cys Asp Ser Lys Arg Gly Tyr Ile Cys Gln Thr Arg Ser 1070 1075 1080
- Asp Pro Ser Leu Thr Asn Pro Pro Ala Thr Ile Gln Thr Asp Gly 1085 1090
- Phe Val Lys Tyr Gly Lys Ser Ser Tyr Ser Leu Met Arg Gln Lys 1100 1105 1110
- Phe Gln Trp His Glu Ala Glu Thr Tyr Cys Lys Leu His Asn Ser 1115 1120 1125
- Leu Ile Ala Ser Ile Leu Asp Pro Tyr Ser Asn Ala Phe Ala Trp 1130 1135 1140
- Leu Gln Met Glu Thr Ser Asn Glu Arg Val Trp Ile Ala Leu Asn 1145 1150 1155
- Ser Asn Leu Thr Asp Asn Gln Tyr Thr Trp Thr Asp Lys Trp Arg 1160 1165 1170

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- Ala Cys Val Tyr Leu Asp Leu Asp Gly Tyr Trp Lys Thr Ala His 1190 1195 1200
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- Leu Leu Thr Thr Lys Ala Asp Thr Arg Lys Met Asp Pro Ser Lys
- Pro Ser Ser Asn Val Ala Gly Val Val Ile Ile Val Ile Leu Leu 1385 1390 1395

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tco Ser 465	Th:	cac His	c tgt s Cys	cco Pro	c tca Ser 470	: Asr	aat Asr	gto Val	att Ile	tca Ser 475	: GI	a aga ı Arç	ı aat J Ası	agt Ser	aca Thr 480	1440
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- Gly Leu Lys Met Gln Arg Thr Val Asn Thr Ile Trp Phe Leu His Leu 50 55 60
- Thr Leu Ala Asp Leu Leu Cys Cys Leu Ser Leu Pro Phe Ser Leu Ala 65 70 75 80
- His Leu Ala Leu Gln Gly Gln Trp Pro Tyr Gly Arg Phe Leu Cys Lys 85 90 95
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Phe Pro Ile Glu Asp His Glu Thr Ser Pro Leu Asp Asn Ser Asp Ala 275 280 285

Phe Leu Ser Thr His Leu Lys Leu Phe Pro Ser Ala Ser Ser Asn Ser 290 295 300

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_		_		-	cgc Arg							_		_		3523
					att Ile 345											3571
	_	_			gta Val	_			_	-			-	-	-	3619
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	_		-		tta Leu									-	-	3715
	_	_		_	tta Leu		_				_					3763
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Ile Val Ile Tyr Ala Tyr Ile Thr Lys Ile Ser Asp Ser Ile Phe Val 100 105 110

Ile Phe Pro Ser Phe Phe Val Val Tyr Val Thr Ile Ser Ile Leu Phe 115 120 125

Ser Tyr Ile Ile Asn Arg Val Leu Lys Lys Ile Ser Thr Pro Tyr Leu 130 135 140

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Thr Phe Ser Leu Phe Phe Phe Tyr Ser Gln Ile Asn Ser Asp Glu Ala 165 170 175

Lys Val Ile Arg Gln Tyr Ser Leu Phe Tyr Trp Tyr His Tyr Ile Leu 180 185 190

Ser Ile Leu Thr Leu Tyr Ser Gln Phe Leu Leu Lys Glu Met Lys Tyr 195 200 205

Lys Arg Asn Gln Glu Glu Ile Glu Thr Tyr Tyr Glu Tyr Thr Leu Lys 210 215 220

Ile Glu Ala Ile Asn Asn Glu Met Arg Lys Phe Arg His Asp Tyr Val 225 230 235 240

Asn Ile Leu Thr Thr Leu Ser Glu Tyr Ile Arg Glu Asp Asp Met Pro 245 250 255

Gly Leu Arg Asp Tyr Phe Asn Lys Asn Ile Val Pro Met Lys Asp Asn 260 265 270

Leu Gln Met Asn Ala Ile Lys Leu Asn Gly Ile Glu Asn Leu Lys Val 275 280 285

Arg Glu Ile Lys Gly Leu Ile Thr Ala Lys Ile Leu Arg Ala Gln Glu 290 295 300

Met Asn Ile Pro Ile Ser Ile Glu Ile Pro Asp Glu Val Ser Ser Ile 305 310 315 320

Asn Leu Asn Met Ile Asp Leu Ser Arg Ser Ile Gly Ile Ile Leu Asp 325 330 335

Asn Ala Ile Glu Ala Ser Thr Glu Ile Asp Asp Pro Ile Ile Arg Val $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$

Ala Phe Ile Glu Ser Glu Asn Ser Val Thr Phe Ile Val Met Asn Lys 355 360 365

Cys Ala Asp Asp Ile Pro Arg Ile His Glu Leu Phe Gln Glu Ser Phe 370 375 380

Ser Thr Lys Gly Glu Gly Arg Gly Leu Gly Leu Ser Thr Leu Lys Glu 385 390 395 400

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Asp Ile Asn Gly Ile Lys Leu Gly Ser Glu Ile Arg Lys His Asp Pro 65 70 75 80

Val Gly Asn Ile Ile Phe Val Thr Ser His Ser Glu Leu Thr Tyr Leu 85 90 95

Thr Phe Val Tyr Lys Val Ala Ala Met Asp Phe Ile Phe Lys Asp Asp 100 105 110

Pro Ala Glu Leu Arg Thr Arg Ile Ile Asp Cys Leu Glu Thr Ala His 120 115 Thr Arg Leu Gln Leu Leu Ser Lys Asp Asn Ser Val Glu Thr İle Glu 135 Leu Lys Arg Gly Ser Asn Ser Val Tyr Val Gln Tyr Asp Asp Ile Met 150 155 Phe Phe Glu Ser Ser Thr Lys Ser His Arg Leu Ile Ala His Leu Asp 170 Asn Arg Gln Ile Glu Phe Tyr Gly Asn Leu Lys Glu Leu Ser Gln Leu 180 185 Asp Asp Arg Phe Phe Arg Cys His Asn Ser Phe Val Val Asn Arg His 200 195 Asn Ile Glu Ser Ile Asp Ser Lys Glu Arg Ile Val Tyr Phe Lys Asn Lys Glu His Cys Tyr Ala Ser Val Arg Asn Val Lys Lys Ile 225 230 235 <210> 105 <211> 7499 <212> DNA <213> Escherichia coli <220> <221> CDS <222> (444)..(1475) <223> <220> <221> CDS <222> (1478)..(2590) <223> <220> <221> CDS <222> (2592)..(3386) <223> <220> <221> CDS <222> (3426)..(4361) <223> <220> <221> CDS <222> (4375)..(5481) <223> <220> <221> CDS <222> (5484)..(6503) <223> <220> <221> CDS <222> (6543)..(7487) <223> <400> 105 60 atagtaaata gctgacttat ggatgtgctg gggattctat gtatttagct gtggctttac cattactttt cccqtttttq acttaaataq cttcaqtttq qtctqatctq ccqctacatc 120 ttcatttttt ttgtattttt atgcgattca ttgaaactcg gccccatttt caaatctaca 180 taggccgtac tgacattatc gaaatgctat tttttatcta tttgattttt atgattaaag 240 tatattttgt gtataaaaat cattcgggtc ggattgctgc gaaagaaatg atacactagc 300 acgtcaaagt aagtgcgtta tcagtattca ggtagctgtt gagcctgggg cggtagcgtg 360

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act cct gtc atc agt acg ctc aag cag aat tat cct gat gca aaa atc Thr Pro Val Ile Ser Thr Leu Lys Gln Asn Tyr Pro Asp Ala Lys Ile 15 20 25	521
gat atg ctg ctt tat cag gac acc atc cct att ttg tct gaa aac ccg Asp Met Leu Tyr Gln Asp Thr Ile Pro Ile Leu Ser Glu Asn Pro 30 35 40	569
gaa att aat gcg ctc tat ggg ata agc aat aaa ggt gcg gga act ttc Glu Ile Asn Ala Leu Tyr Gly Ile Ser Asn Lys Gly Ala Gly Thr Phe 45 50 55	617
gat aaa att aaa aat gtg ctt tcg ttg ata aaa act ctg cgt gcg aat Asp Lys Ile Lys Asn Val Leu Ser Leu Ile Lys Thr Leu Arg Ala Asn 60 65 70	665
aat tat gac ctg gtc att aat ctt acg gat cag tgg atg gtg gcg ctg Asn Tyr Asp Leu Val Ile Asn Leu Thr Asp Gln Trp Met Val Ala Leu 75 80 85 90	713
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ata cac ggt aca cat att gtt gag cgt aat tta tcg gtc ctt gag cca Ile His Gly Thr His Ile Val Glu Arg Asn Leu Ser Val Leu Glu Pro 125 130 135	857
tta ggt att acc gat ttc tac acc gac aca aca atg agt tac gcc gaa Leu Gly Ile Thr Asp Phe Tyr Thr Asp Thr Thr Met Ser Tyr Ala Glu 140 145 150	905
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cat His 650	tat Tyr	att Ile	gcg Ala	gat Asp	gcc Ala 655	aat Asn	tgt Cys	gga Gly	acg Thr	gtc Val 660	atc Ile	gct Ala	gaa Glu	cct Pro	ttc Phe 665	:	2440
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						tta Leu										3161
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	_	Leu I				_	yr Le	_			-	yr Ly	-	ita aaa Ile Lys	3506
	ata Ile	aca Thr 1010					tat Tyr 1015						aaa Lys		3551
		ata Ile 1025		-	-		gca Ala 1030		_		_	tta Leu 1035	ttc Phe		3596
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- ·	-	ttt Phe 1085					gcg Ala 1090					att Ile 1095			3776
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			Phe				atg Met 1120								3866
			Lys				cag Gln 1135							ggt Gly	3911
	gat Asp		Gly				ata Ile 1150					gcc Ala 1155		aga Arg	3956
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		gca Ala 1220					aaa Lys 1225							ata Ile	4181
		ttt Phe 1235					tat Tyr 1240					ttt Phe 1245	ttt Phe	ata Ile	4226
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	a agt n Ser 133	Ğlu						_		_		Met	_	_	4545
	a tgg y Trp 135			ggt Gly			Tyr		_		ttt Phe 136	Ser	aat Asn		4590
_	t tta g Leu 136	Ğİy		ttg Leu	_		Ala				cat His 137	Ala	cta Leu	_	4635
aa	a tgg	ctt	cag	gag	tac	caa	cca	gat	atc	gtt	att	tgt	att	gat	4680

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	gtc Val 1470					gag Glu 1475									4950
						ttt Phe 1490									4995
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						gaa Glu 1685									5582
						cta Leu 1700									5627
	ata Ile 1710					gaa Glu 1715		_	_	_	_				5672
		_				gat Asp 1730	_	-	_	_			-	_	5717
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						ttg Leu 1760							act Thr		5807
						gtt Val 1775					ttt Phe 1780		aat Asn	_	5852
	cct Pro 1785					ctg Leu 1790									5897
	att Ile 1800					aat Asn 1805									5942
	atg Met 1815					ggg Gly 1820						gaa Glu			5987
gcc	cat	tcg	tta	ggt	gtt	gcg	gga	att	gct	aaa	ggt	tac	ttt	aac	6032

Ala	His 1830	Ser	Leu	Gly	Val	Ala 1835		Ile	Ala	Lys	Gly 1840		Phe	Asn	
Ser						aat Asn 1850	Thr				gcg Ala 1855	Ala			6077
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	aaa Lys 1875					gat Asp 1880		-			aat Asn 1885	Meť	ttg Leu	_	6167
						gct Ala 1895	Āsp					Thr	cag Gln		6212
						aaa Lys 1910							gta Val		6257
	gat Asp 1920					cat His 1925					acc Thr 1930	Lys	ccc Pro		6302
						tat Tyr 1940	Pro				gcc Ala 1945	Phe	atg Met		6347
	aaa Lys 1950					tgg Trp 1955	Lys					Leu	aaa Lys		6392
	aat Asn 1965					aga Arg 1970		_	_		cat His 1975	-	ctt Leu		6437
_	cat His 1980					gga Gly 1985	Phe					Phe			6482
	gaa Glu 1995	_		aag Lys		taa	aacto	gggaa	aa ci	tataa	aagta	atga	atata	aag	6533
ggta	igcati		Ĩ Ās			tt cc ne Pr		a I	_	-			s Vá	tt aaa al Lys	6581
_	Trp A	_				gct Ala					Ser	_	-		6626
	gtt o Val i	_				gat Asp					Āsp			ggt Gly	6671
gtt	tcc a	atc	aca	tca	att	gtç	cta	aat	aat	cga	cat	att	aat	ctt	6716

Val	Ser	Ile	Thr	Ser	Ile	Val	Leu	Asn	Asn	Arg	His	Ile	Asn	Leu	
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		gca Ala 2075					caa Gln 2080					atc Ile 2085			6806
	-				_	_	ctc Leu 2095					tgt Cys 2100		-	6851
_			-	-	_		ttc Phe 2110	_			_				6896
-			_		_	-	ttg Leu 2125				-		-	-	6941
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		gtt Val 2150	-	_	_	_	aaa Lys 2155					atg Met 2160	caa Gln	-	7031
							gat Asp 2170					ggg Gly 2175			7076
							tta Leu 2185	_				tgg Trp 2190	-	-	7121
		cta Leu 2195		-			ttg Leu 2200					agt Ser 2205		gat Asp	7166
							cag G1n 2215					gtc Val 2220		ctg Leu	7211
							cct Pro 2230					aca Thr 2235		tat Tyr	7256
		aaa Lys 2240	_	_			gac Asp 2245					aac Asn 2250		aaa Lys	7301
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	gat Asp	gcg Ala 2300	Lys			att Ile	_	ı Ph			_	p I		aac Asn	_	7481
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Asp	Thr	Ile 35	Pro	Ile	Leu	Ser	Glu 40	Asn	Pro	Glu	Ile	Asn 45	Ala	Le:	ı Tyr	
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Asn	Leu	Thr	Asp	Gln 85	Trp	Met	Val	Ala	Leu 90	Leu	Val	Arg	Cys	95	ı Pro	
Ala	Arg	Met	Lys 100	Ile	Ser	Gln	Leu	Tyr 105	Gly	His	Arg	Gln	His 110		y Ile	
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Tyr 145	Thr	Asp	Thr	Thr	Met 150	Ser	Туr	Ala	Glu	Asp 155	Cys	Trp	Lys	s Ly	s Met 160	
Arg	Arg	Glu	Leu	Asp 165	Ala	Leu	Gly	Val	Lys 170	Asp	His	Tyr	Va.	l Va 17	l Ile 5	

Gln Pro Thr Ala Arg Gln Ile Phe Lys Cys Trp Asp Asn Asp Lys Phe 180 185 190

Ser Lys Val Ile Asp Ala Leu Gln Gln Arg Gly Tyr Gln Val Val Leu 195 200 205

Thr Cys Gly Pro Ser Ala Asp Asp Leu Ala Cys Val Asp Glu Ile Ala 210 215 220

Arg Gly Cys Glu Thr Lys Pro Ile Thr Gly Leu Ala Gly Lys Thr Arg 225 230 235 240

Phe Pro Glu Leu Gly Ala Leu Ile Asp His Ala Val Leu Phe Ile Gly 245 250 255

Val Asp Ser Ala Pro Gly His Ile Ala Ala Ala Val Lys Thr Pro Val 260 265 270

Ile Ser Leu Phe Gly Ala Thr Asp His Val Phe Trp Arg Pro Trp Thr 275 280 285

Glu Asn Ile Ile Gln Phe Trp Ala Gly Asn Tyr Gln Lys Met Pro Thr 290 295 300

Arg His Glu Leu Asp Arg Asn Lys Lys Tyr Leu Ser Val Ile Pro Ala 305 310 315 320

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- Glu Leu Ile Gln Val Pro Val Lys Ser His Thr Asn His Gly Arg Asn 50 60
- Ala Glu Tyr Tyr Ala Trp Val Gln Asn His Leu Lys Glu His Pro Ala 65 70 75 80
- Asp Arg Val Val Gly Phe Asn Lys Met Pro Gly Leu Asp Val Tyr Phe 85 90 95
- Ala Ala Asp Val Cys Tyr Ala Glu Lys Val Ala Gln Glu Lys Gly Phe 100 105 110
- Leu Tyr Arg Leu Thr Ser Arg Tyr Arg His Tyr Ala Ala Phe Glu Arg 115 120 125
- Ala Thr Phe Glu Gln Gly Lys Ser Thr Lys Leu Met Met Leu Thr Asp 130 135 140
- Lys Gln Ile Ala Asp Phe Gln Lys His Tyr Gln Thr Glu Pro Glu Arg 145 150 155 160
- Phe Gln Ile Leu Pro Pro Gly Ile Tyr Pro Asp Arg Lys Tyr Ser Glu 165 170 175
- Gln Ile Pro Asn Ser Arg Glu Ile Tyr Arg Gln Lys Asn Gly Ile Lys 180 185 190
- Glu Gln Gln Asn Leu Leu Gln Val Gly Ser Asp Phe Gly Arg Lys 195 200 205
- Gly Val Asp Arg Ser Ile Glu Ala Leu Ala Ser Leu Pro Glu Ser Leu 210 215 .220
- Arg His Asn Thr Leu Leu Phe Val Val Gly Gln Asp Lys Pro Arg Lys 225 230 235 240
- Phe Glu Ala Leu Ala Glu Lys Leu Gly Val Arg Ser Asn Val His Phe 245 250 255
- Phe Ser Gly Arg Asn Asp Val Ser Glu Leu Met Ala Ala Ala Asp Leu 260 265 270
- Leu Leu His Pro Ala Tyr Gln Glu Ala Ala Gly Ile Val Leu Leu Glu 275 280 285

Ala Ile Thr Ala Gly Leu Pro Val Leu Thr Thr Ala Val Cys Gly Tyr 290 295 300

Ala His Tyr Ile Ala Asp Ala Asn Cys Gly Thr Val Ile Ala Glu Pro 305 310 315 320

Phe Ser Gln Glu Gln Leu Asn Glu Val Leu Arg Lys Ala Leu Thr Gln 325 330 335

Ser Pro Leu Arg Met Ala Trp Ala Glu Asn Ala Arg His Tyr Ala Asp 340 345 350

Thr Gln Asp Leu Tyr Ser Leu Pro Glu Lys Ala Ala Asp Ile Ile Thr . 355 360 365

Gly Gly Leu 370

Ala Phe Ala Glu Val Lys Lys Leu As
n Gly Glu Val Phe Arg Glu Leu 20 25 30

Glu Thr Arg Arg Thr Leu Arg Phe Glu Leu Ser Gly Lys Ser Tyr Phe 35 40 45

Leu Lys Trp His Lys Gly Thr Thr Leu Lys Glu Ile Ile Lys Asn Leu 50 60

Leu Ser Leu Arg Met Pro Val Leu Gly Ala Asp Arg Glu Trp His Ala 65 70 75 80

Ile His Arg Leu Ser Asp Val Gly Val Asp Thr Met Lys Gly Ile Gly 85 90 95

Phe Gly Glu Lys Gly Leu Asn Pro Leu Thr Arg Ala Ser Phe Ile Ile 100 105 110

Thr Glu Asp Leu Thr Pro Thr Ile Ser Leu Glu Asp Tyr Cys Ala Asp 115 120 125

Trp Ala Val Asn Pro Pro Asp Ile Arg Val Lys Arg Met Leu Ile Ala 130 135 Arg Asp Cys Tyr Ile Cys His Phe Leu Leu His Leu Pro Phe Thr Gly 165 170 175

Arg Glu Asp Glu Leu Lys Ile Ser Val Ile Asp Leu His Arg Ala Gln 180 185 190

Ile Arg Ala Lys Val Pro Arg Arg Trp Arg Asp Lys Asp Leu Ile Gly 195 200 205

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Arg Phe Met Lys Val Tyr Phe Gly Met Pro Leu Arg Lys Ile Leu Ser 225 230 235 240

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Lys Lys Tyr Asn Trp Ile Asn Lys Lys Ile Ser Lys Ile Gly Met Asp 35 40 45

Asn Ala Trp Leu Gly Arg Leu Phe Ile Lys His Tyr Leu Asn Asp Ile 50 55 60

Lys Lys Asp Asp Ile Ile Ile Ile Asn Asp Ser Val Val Asn Lys Gly 65 70 75 80

Ile Asn Lys Gln Ile Leu Lys Asn Ile Asn Cys His Lys Val Leu Leu 85 90 95

Phe Asp Ile Ile Tyr Asp Phe Glu His Arg Phe Ile Gly Asn Glu Lys 115 120 125

Ile Lys Ala Ile Glu Gln Phe Phe Pro Ile Gly Met Asp Glu Ile Arg 130 135 140

Asn Tyr Ser Leu Ser Asp Lys Asn Asn Ser Gln Pro Ile Cys Phe Phe 145 150. 155 160

Leu Gly Arg Asp Lys Gly Arg Leu Gln Ile Ile Asn Glu Leu Ala Glu 165 170 175

Arg Leu Thr Thr Leu Gly Cys Lys Leu Asp Phe Asn Val Val Lys Asp 180 185 190

Lys Thr Ser Ser Thr Thr Ser Lys Tyr Leu Ile Glu Lys Gln Ile Ser 195 200 205

Tyr Glu Glu Asn Ile Arg Arg Thr Leu Asn Ala Asn Ile Ile Val Asp 210 215 220

Ile Thr Lys Glu Asn Gln Ser Gly Trp Thr Leu Arg Ile Leu Glu Ala 225 230 235 240

Leu Phe Phe Asn Lys Lys Leu Ile Thr Asn Asn Ile Asn Val Phe Gly 245 250 255

Ser Glu Ile Tyr Ser Glu Ser Arg Phe Phe Ile Ile Gly His Asp Asp 260 265 270

Trp Asp Lys Leu Glu Tyr Phe Ile Asn Ser Ser Val Lys Pro Met Asp 275 280 285

Tyr Asp Ser Leu Tyr Lys Phe Ser Pro Asp Lys Met Met Ser Thr Ile 290 295 300

Val Ser Asp Phe Ile Asp Lys

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- Gly Glu Ala Val Ser Gly Phe Gly Gly Met Glu Thr Val Ile Arg Asp 20 25 30
- Val Ile Thr Thr Phe Arg Gln Gln His Ile Gln Ser Glu Met Phe 35 40 45
- Phe Cys Arg Asn Asp Lys Met Asp Lys Gly Trp Leu Glu Gly Ile Lys 50 55 60
- Tyr Ser Cys Ser Phe Ser Asn Ile Arg Leu Gly Phe Leu Arg Arg Ala 70 75 80
- Lys His Ile His Ala Leu Ser Lys Trp Leu Gln Glu Tyr Gln Pro Asp 85 90 95
- Ile Val Ile Cys Ile Asp Val Ile Ser Cys Leu Phe Ala Ala Lys Ala 100 105 110
- Arg Lys Ser Gly Ile Asp Met Pro Val Phe Ser Trp Pro His Phe 115 120 125
- Ser Leu Asp His Lys Lys His Ala Glu Tyr Ile Thr Cys Ala Asp Tyr 130 135 140
- His Leu Ala Ile Ser Ser Gly Ile Lys Gln Gln Met Ile Asn Arg Gly 145 150 155 160
- Val Ala Glu Ser Thr Ile Asn Val Ile Phe Asn Pro Val Glu Thr Lys 165 170 175
- Asp Ser Val Ile Pro Ala Pro Glu Glu Gly Glu Thr Ala Thr Phe Ile 180 185 190
- Tyr Val Gly Arg Met Lys Phe Glu Gly Gln Lys Arg Val Lys Asp Leu 195 200 205
- Leu Asp Gly Leu Ser Gln Ala Lys Gly Asn Trp Lys Leu His Val Leu 210 215 220
- Gly Asp Gly Ser Asp Phe Glu Lys Cys Gln Ala Tyr Gly Arg Glu Leu 225 230 230 235 240
- Asn Ile Asp Asp Arg Ile Val Trp Tyr Gly Trp Gln Gln Tyr Pro Trp 245 250 255

- Glu Leu Val Gln Gln Asp Ile Glu Lys Val Ser Ala Leu Leu Thr 260 265 270
- Ser Ser Phe Glu Gly Phe Pro Met Thr Leu Leu Glu Ala Leu Ser Trp 275 280 285
- Gly Ile Pro Cys Ile Ser Ala Asp Cys Val Ser Gly Pro Ala Asp Ile 290 295 300
- Ile Gln Pro Asp Val Asn Gly His Leu Tyr Gln Pro Gly Asp Ile Ala 305 310 315 320
- Gly Phe Val Thr Leu Leu Asn Lys Tyr Ile Ala Gly Glu Ile His Ile 325 330 335
- Glu His Glu Lys Ile Pro Ala Ser Ile Asp Glu Phe Tyr Gln Ser Lys 340 345 350
- Tyr Tyr Asp Arg Leu His Lys Val Ile Ile Ser Ala Ile Ser Arg Arg 355 360 365

Lys

- Asp Tyr Asp His Lys Val Glu Thr Glu Asn Leu Cys Leu Asp Ile Ala 20 25 30
- Tyr Gly Thr Asp Lys Asn Phe Leu Phe Gly Cys Gly Ile Ser Ile Ala 35 40 45
- Ser Ile Leu Lys Tyr Asn Glu Gly Ser Arg Leu Cys Phe His Ile Phe 50 55 60
- Thr Asp Tyr Phe Gly Asp Asp Asp Arg Lys Tyr Phe Asp Ala Leu Ala 65 70 75 80
- Leu Gln Tyr Lys Thr Arg Ile Lys Ile Tyr Leu Ile Asn Gly Asp Arg

- Leu Arg Ser Leu Pro Ser Thr Lys Asn Trp Thr His Ala Ile Tyr Phe 100 105 110
- Arg Phe Val Ile Ala Asp Tyr Phe Ile Asn Lys Ala Pro Lys Val Leu 115 120 125
- Tyr Leu Asp Ala Asp Ile Ile Cys Gln Gly Thr Ile Glu Pro Leu Ile 130 135 140
- Asn Phe Ser Phe Pro Asp Asp Lys Val Ala Met Val Val Thr Glu Gly 145 150 155 160
- Gln Ala Asp Trp Trp Glu Lys Arg Ala His Ser Leu Gly Val Ala Gly 165 170 175
- Ile Ala Lys Gly Tyr Phe Asn Ser Gly Phe Leu Leu Ile Asn Thr Ala 180 185 190
- Gln Trp Ala Ala Gln Gln Val Ser Ala Arg Ala Ile Ala Met Leu Asn 195 200 205
- Glu Pro Glu Ile Ile Lys Lys Ile Thr His Pro Asp Gln Asp Val Leu 210 215 220
- Asn Met Leu Leu Ala Asp Lys Leu Ile Phe Ala Asp Ile Lys Tyr Asn 225 230 235 240
- Thr Gln Phe Ser Leu Asn Tyr Gln Leu Lys Glu Ser Phe Ile Asn Pro 245 250 255
- Val Thr Asn Asp Thr Ile Phe Ile His Tyr Ile Gly Pro Thr Lys Pro 260 265 270
- Trp His Asp Trp Ala Trp Asp Tyr Pro Val Ser Gln Ala Phe Met Glu 275 280 280
- Ala Lys Asn Ala Ser Pro Trp Lys Asn Thr Ala Leu Leu Lys Pro Asn 290 295 300
- Asn Ser Asn Gln Leu Arg Tyr Ser Ala Lys His Met Leu Lys Lys His 305 310 315 320
- Arg Tyr Leu Lys Gly Phe Ser Asn Tyr Leu Phe Tyr Phe Ile Glu Lys 325 330 335

Ile Lys His

Phe Arg Leu Ala Asn Ile Asn Thr Ser Glu Cys Leu Asn Val Ala Tyr 20 25 30

Gly Val Asp Ala Asn Tyr Leu Asp Gly Val Gly Val Ser Ile Thr Ser 35 40 45

Ile Val Leu Asn Asn Arg His Ile Asn Leu Asp Phe Tyr Ile Ile Ala 50 55 60

Asp Val Tyr Asn Asp Gly Phe Phe Gln Lys Ile Ala Lys Leu Ala Glu 65 70 75 80

Gln Asn Gln Leu Arg Ile Thr Leu Tyr Arg Ile Asn Thr Asp Lys Leu 85 90 95

Gln Cys Leu Pro Cys Thr Gln Val Trp Ser Arg Ala Met Tyr Phe Arg 100 105 110

Leu Phe Ala Phe Gln Leu Leu Gly Leu Thr Leu Asp Arg Leu Leu Tyr 115 120 125

Leu Asp Ala Asp Val Val Cys Lys Gly Asp Ile Ser Gln Leu Leu His 130 135 140

Leu Gly Leu Asn Gly Ala Val Ala Ala Val Val Lys Asp Val Glu Pro 145 150 155 160

Met Gln Glu Lys Ala Val Ser Arg Leu Ser Asp Pro Glu Leu Leu Gly 165 170 175

Gln Tyr Phe Asn Ser Gly Val Val Tyr Leu Asp Leu Lys Lys Trp Ala 180 185 190

Asp Ala Lys Leu Thr Glu Lys Ala Leu Ser Ile Leu Met Ser Lys Asp 195 200 205

Asn Val Tyr Lys Tyr Pro Asp Gln Asp Val Met Asn Val Leu Leu Lys 210 215 220

Gly 225	Met	Thr	Leu	Phe	Leu 230	Pro	Arg	Glu	Tyr	Asn 235	Thr	Ile	Tyr	Thr	Ile 240	
Lys	Ser	Glu	Leu	Lys 245	Asp	Lys	Thr	His	Gln 250	Asn	Tyr	Lys	Lys	Leu 255	Ile	
Thr	Glu	Ser	Thr 260	Leu	Leu	Ile	His	Tyr 265	Thr	Gly	Ala	Thr	Lys 270	Pro	Trp	
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cag Gln	ggc Gly	cgc Arg	atg Met 580	cct Pro	cca Pro	gcc Ala	gag Glu	tca Ser 585	gct Ala	cgg Arg	tgc Cys	cga Arg	cat His 590	cca Pro	cga Arg	1776
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cgg Arg 785	atg Met	gat Asp	gaa Glu	ccc Pro	gtc Val 790	ggt Gly	gag Glu	ttg Leu	ctg Leu	gac Asp 795	gct Ala	ttc Phe	gag Glu	caa Gln	gcc Ala 800	2400
gca Ala	atc Ile	gat Asp	gaa Glu	gtg Val 805	ctc Leu	ggc Gly	gcc Ala	ggt Gly	gtc Val 810	gag Glu	ccg Pro	aag Lys	gat Asp	gtc Val 815	gcg Ala	2448
tcc Ser	ggc Gly	cgg Arg	ctg Leu 820	Gly ggc	cgg Arg	gcc Ala	gac Asp	gtg Val 825	gcc Ala	gga Gly	ccg Pro	ttg Leu	gct Ala 830	gtc Val	gtc Val	2496
											acc Thr					2544

835		840	845	
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		Ser Ser Thr G	gc gcc cgg ctg cag a ly Ala Arg Leu Gln 1 75	
			gt ctc ggg cac ctg o rg Leu Gly His Leu G 895	
			gt cga tgg cgg cac (rg Arg Trp Arg His 1 910	
	2 22 2		tg cgc acg gtg ctg (let Arg Thr Val Leu <i>l</i> 925	-
J J	, , ,		gc tgc ggt ggc caa (ys Cys Gly Gly Gln 1 940	
, ,,	, ,,	Leu Ala Pro A	ga geg tgt tge ega (rg Ala Cys Cys Arg 1 55	
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	2 2 2 2		cca gcg gtt ttc gcg o Pro Ala Val Phe Ala 7 990	
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	g gtg cat ctg ga 1 Val His Leu As 10	o Thr Arg Pro	g ege gtg gte ggt ea Arg Val Val Gly G 1020	
		n Leu Thr Val	acg caa cgg ctg c Thr Gln Arg Leu P 1035	
-	g ata cgg aca tg o Ile Arg Thr Tr 10	p Ala Ala Ser	g tgc cgg tct cgg tc Cys Arg Ser Arg So 1050	-
	c gca tgg cgc cg r Ala Trp Arg Ar 10	g Asp Arg His	c tot oga gga gog a s Ser Arg Gly Ala I 1065	
	t ggg teg eac eg o Gly Ser His Ar		g ctg gac - cgg cgc g n Leu Asp - Arg Arg G	-

	1070					1075					1080				
ccg (Pro \						gaa Glu 1090									3294
_		_	_	_		acc Thr 1105	~ ~	_	_	_	_	_	_		3339
gcg (Ala V						cac His 1120									3384
						gga Gly 1135									3429
gct (Ala						aca Thr 1150									3474
_						ctg Leu 1165									3519
Gly I						acg Thr 1180									3564
cgg Arg						gag Glu 1195					gcc Ala 1200		_	_	3609
Gly		_			_	tcg Ser 1210					ctg Leu 1215		gca Ala		3654
						ccc Pro 1225									3699
						gcc Ala 1240									3744
Gly	aca Thr 1250	ccg Pro	cgg Arg	aca Thr	agt Ser	tca Ser 1255	ccc Pro	gcg Ala	aca Thr	ccc Pro	tgg Trp 1260	Ala	tct Ser	cgg Arg	3789
	tgc Cys 1265					aca Thr 1270					tça Ser 1275		cca Pro		3834
vаĺ	_					gac Asp 1285					ctg Leu 1290	_	cag Gln		3879
acc Thr						acg Thr									3924

	1295					1300					1305				
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											ggc Gly 1365				4104
											gac Asp 1380				4149
gcc Ala	gac Asp 1385	gtg Val	ccc Pro	gcg Ala	ttc Phe	gtc Val 1390	gcc Ala	Gly ggg	atc Ile	gcg Ala	gag Glu 1395	agc Ser	acc Thr		4194
											ggc Gly 1410		caa Gln		4239
											ctc Leu 1425				4284
	gag Glu 1430		_	_		ctc Leu 1435					cgg Arg 1440		ttc Phe		4329
_	gtg Val 1445										cga Arg 1455				4374
											cgg Arg 1470				4419
	acg Thr 1475										cat His 1485		caa Gln		4464
	gcc Ala 1490					ccc Pro 1495					tca Ser 1500		agg Arg		4509
	ggg Gly 1505					gcc Ala 1510					gag Glu 1515		ctc Leu		4554
	tac Tyr					cgc Arg					gat Asp				4599

Ala Leu Ser Ala Lys Met Arg 1675 tcc atc gag tcc atc acc gac ggt gcg tcg tcg cgg cgc aac cag 5094 Ser Ile Glu Ser Ile Thr Asp 1690 ctg ctg tg gtg gac ctg ggc tcc gag ctg aac ctc ggt Arg Arg Asn Gln 1695 ctg ctg teu Leu Val Asp Leu Gly Ser Glu Leu Asn Leu Gly Ala Ile Glu 1700 cgg cgc cgc cga atc gga cct ggc cgg tct gcg cgc att gaa 5139 Arg Arg Arg Ile Gly Pro Gly Arg Ser Ala Leu Thr Gly Asp 1715 caa act ggc gcg cac cta caa cgt tac ggc cca gtg ctt tcc gac Gln Thr Gly Ala His Leu Gln Arg Tyr Gly Pro Val Leu Ser Asp 1735														-		
Sist Sist		1520					1525					1530				
Leu Asp		Val					Gly					Gln				4644
Giy Leu Giy Val Glu Arg Phe Val Glu Ile Gly Val Lys Ser Ser ccg acg gtg gcg ggg tt cca caa cac		Asp					Ser					Gly				4689
The Thr 1580 Val Ala Gly Ser Cys 1585 Fis Gln His Pro Gln Thr Ala Arg 1590 Fro Gln His Ser Glu 1590 Fro Gln His Ser Glu 1600 Fro Glu 1600		Leu					Phe					Val				4734
The Arg 1595 For Gln His Ser Glu 1600 For Leu Asn Ala Glu 1605 For Arg Arg Arg Arg Arg Arg Cys Cys Ser Pro Pro Thr 1615 For Thr Arg Ser Arg Arg 1620 For Arg Arg 1610 For Thr Arg Ser Arg Arg 1620 For Arg 1620 For Ar		Thr	gtg Val	gcg Ala	ggg Gly	tct Ser	Cys	cac	caa Gln	cac His	cct Pro	Gln				4779
Arg Cys 1610 aag acg acg cgg tcg cgg aat cgc ccg ccg cgc cgg acg tcc ccc ccc tcg cgc ccg ccg cgc cgc c		Arg					Glu					Glu				4824
Lys Thr 1625 Ser Arg Ser Arg Asn 1630 Arg Pro Arg Arg Thr 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1635 Ser Ser Arg 1645 Ser Ser Arg 1640 Ser Arg Arg Arg Arg Arg Arg Arg Arg Arg Ar		Cys					Thr					Arg				4869
Lys Pro 1640 Pro Pro Ser Arg Arg 1645 Ccg acg at tegg ttt teg acg ccg ccg atg cca ccg tgc gtg atc cys Val Ile 1655 Gcg ctc tcg gcc aag atg ccc atg atg cag atg atg cca cgc tgc Val Ile 1655 Gcg ctc tcg gcc aag atg ccc atg atg cag atg atg atg atg atg atg atg atg atg a	_	Thr	_		_		Asn	_	_	_		Thr	_			4914
Pro Thr Ile Trp Phe Ser Thr 1660 Pro Met Pro Arg Cys Val Ile 1655 gcg ctc tcg gcc aag atg cgc atc gac cag atc gaa gaa ctc gac Ser Ala Lys Met Arg 1675 tcc atc gag tcc atc acc gac ggt gcg tcg tcg cgg cgc aac cag Arg Arg Arg Arg Ile Thr Asp 1690 ctg ctg gtg gac ctg ggc tcc gag ctg aac ctc ggt gcc att gaa Arg Arg Arg Arg Ile Gly Pro 1705 cgg cgc cgc cga atc gga cct ggc cgc cgg ctc ggt gcc att gaa Arg Arg Arg Arg Ile Gly Pro 1715 caa act ggc gcg cac cta caa cgt cgc acc gtc ggc cca gtg ctt tcc gac Gly Arg Ser Ala Leu Thr Gly Asp 1735 caa act ggc gcg cac cta caa cgt tcg cac gtc ctc ggc ctc acc ggc cac cta caa cgt fly Arg Ser Ala Leu Thr Gly Asp 1735 gcc atc aac gac cac gtt cgc acc gtc ctc ggc ctc gtg ctc fly Arg Ser Ala Leu Ser Asp 1730 gcc atc aac gac cac gtt cgc acc gtc ctc gga ccg ftcg ggc ctc fly Arg Ser Ala Leu Ser Asp 1740 gcc atc aac gac cac gtt cgc acc gtc ctc gga ccg ftcg ggc aag 5274		Pro					Arg					Arg				4959
Ala Leu Ser Ala Lys Met Arg 1675 tcc atc gag tcc atc acc gac ggt gcg tcg tcg cgg cgc aac cag 5094 Ser Ile Glu Ser Ile Thr Asp 1690 ctg ctg gtg gac ctg ggc tcc gag ctg aac ctc ggt gcc att gaa Ala Ile Glu 1700 cgg cgc cgc cga atc gga cct ggc cgg tcd ggc tcd ggc cgc att gaa Ala Ile Glu 1700 cgg cgc cgc cga atc gga cct ggc cgg tcd ggc ctc ggc ctc aca ggc cgc atc aca ggc cgc atc aca ggc cgc atc aca ggc cgc cgc cgc atc ggc cgc cgc cgc cgc atc ggc cgc cgc atc aca ggc cgc acc cta caa cgt tac ggc cca gtg ctt tcc gac Thr Gly Ala His Leu Gln 1735 gcc atc aac gac cac gtt cgc acc gac ctc cac gtc ctc ggc ccc tcc ggc cgc atc acc gtc ctc gcc cca gtg ctc tcc gac 5229 gcc atc aac gac cac gtt cgc acc gtc ctc ggc ccc tcc ggc ccc tcc gac 5229 gcc atc aac gac cac gtt cgc acc gtc ctc ggc ccc tcc ggc ccc tcc ggc aac 5274		Thr					Thr					Arg				5004
Ser Ile Glu Ser Ile Thr Asp 1690		Leu					Arg					Glu				5049
Leu Leu Val Asp Leu Gly Ser Glu Leu Asn Leu Gly Ala Ile Glu 1700 1705 1710 cgg cgc cgc cga atc gga cct ggc cgg tct gcg ctc aca ggt gac Arg Arg Arg Ile Gly Pro Gly Arg Ser Ala Leu Thr Gly Asp 1715 1720 1725 caa act ggc gcg cac cta caa cgt tac ggc cca gtg ctt tcc gac Gln Thr Gly Ala His Leu Gln Arg Tyr Gly Pro Val Leu Ser Asp 1730 1735 1740 gcc atc aac gac cac gtt cgc acc gtc ctc gga ccg tcg ggc aag 5274		Ile	gag Glu	tcc Ser	atc Ile	acc Thr	Asp	Gly	gcg Ala	tcg Ser	tcg Ser	Arg	cgc Arg	aac Asn	cag Gln	5094
Arg Arg Arg Ile Gly Pro Gly Arg Ser Ala Leu Thr Gly Asp 1715 1720 1725 caa act ggc gcg cac cta caa cgt tac ggc cca gtg ctt tcc gac Gln Thr Gly Ala His Leu Gln Arg Tyr Gly Pro Val Leu Ser Asp 1730 1735 1740 gcc atc aac gac cac gtt cgc acc gtc ctc gga ccg tcg ggc aag 5274	_	Leu					Ser	Glu				Gly	Ala			5139
Gln Thr Gly Ala His Leu Gln Arg Tyr Gly Pro Val Leu Ser Asp 1730 1735 1740 gcc atc aac gac cac gtt cgc acc gtc ctc gga ccg tcg ggc aag 5274		Arg	_	_			Pro					Leu			_	5184
9		Thr					Gln	_				Val	Leu		-	5229
	_								-			_				5274

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	tca Ser 1880					cgc Arg 1885		gat Asp	tcc Ser	gag Glu	ctg Leu 1890		gac Asp		5679
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-	agt Ser					ccg Pro		-			atc Ile	_	_	cgg Arg	5949

	1970					1975					1980				
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	acg Thr 2105												cga Arg		6354
	gat Asp 2120												ggc Gly		6399
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		_				cgc Arg 2215	_	_	_						6669
						ggc Gly 2230									6714
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						gcg Ala 2290					tgc Cys 2295				6894
						ccg Pro 2305									6939
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	gtc Val 2585					gcg Ala 2590								7794
-	acc Thr 2600					gtc Val 2605	Glu							7839
	gga Gly 2615					tgg Trp 2620					gac Asp			7884
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Arg	Gly	Asp 35	Pro	Gly	Ala		ro S 0	Ser A	rg H	is G	ly Pr 45		g Gl	y His		
Pro	Ala 50	Gly	Arg	His	Gly	Gly H 55	lis S	Ser G	ln G		al Le 50	u Al	a Va	l Glu		

- Ala Leu Lys Ala Gly Gly Ala Arg Asp Val Glu Leu Phe Ala Leu Ala 65 70 75 80
- Gln Leu Ile Gly Ala Ala Gly Thr Leu Val Ala Arg Arg Arg Glu Phe 85 90 95
- Pro Ser Trp Ala Ile Ala Pro Met Val Ser Val Thr Asn Ala Asp Pro 100 105 110
- Glu Arg Ile Gly Arg Leu Leu Asp Glu Phe Ala Gln Asp Val Arg Thr 115 120 125
- Val Leu Pro Pro Val Leu Ser Ile Arg Asn Gly Arg Arg Ala Val 130 135 140
- Ile Thr Gly Thr Pro Glu Gln Leu Ser Arg Phe Glu Leu Tyr Cys Arg 145 150 155 160
- Gln Ile Ser Glu Lys Glu Glu Ala Asp Arg Lys Asn Lys Val Arg Gly 165 $$ 170 $$ 175
- Gly Asp Val Phe Ser Pro Val Phe Glu Pro Val Gln Val Gly 180 185 190
- Phe His Thr Pro Arg Leu Ser Asp Gly Ile Asp Ile Val Ala Gly Trp 195 200 205
- Ala Ile Leu Ile Arg Lys Val Asp Trp Val Asp Glu Ile Thr Arg Val 225 230 235 240
- His Arg Ala Gly Ala Arg Trp Ile Leu Asp Leu Gly Pro Gly Asp Ile 245 250 255
- Leu Thr Arg Leu Thr Ala Pro Val Ile Arg Gly Leu Gly Ile Gly Ile 260 265 270
- Val Pro Ala Arg Thr Arg Gly Gly Gln Arg Asn Leu Phe Thr Val Gly 275 280 285
- Ala Thr Pro Glu Val Ala Arg Ala Trp Ser Ser Tyr Ala Pro Thr Val 290 295 300

- Val Arg Leu Pro Asp Gly Arg Val Lys Leu Ser Thr Lys Phe Thr Arg 305 310 315 320
- Val Asp Ala Lys Ile Val Ala Ala Ala Ala Asn Gly Arg His Trp Ala 340 345 350
- Glu Leu Ala Ala Arg Gly Arg Ser Pro Lys Arg Ser Ser Val Thr Ala 355 360 365
- Ser Asn Lys Trp Pro Ala Cys Ser Ser Arg Ala Ala Pro Ile Ser Ser 370 375 380
- Thr Arg Cys Ser Ser Ile Pro Thr Cys Glu Ala Ser Gly Gly Arg Gln 385 390 395 400
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- Arg Arg Thr Gly Arg His Arg His Gln Pro Arg Arg Val Gln Thr Arg 435 440 445
- Asp His Arg Ala Asp Pro Leu Gly Asp Ser His Arg His Arg Gly Ala 450 455 460
- His Gln Ala Gly Asp His Ala Arg Arg Gly Pro Gly Ala Pro Ala Gly 465 470 475 480
- Thr Ile Pro Gly Arg Ile Ser His Leu Leu Leu Ala Thr Tyr Ser Ala 485 490 495
- Asp Arg Ala Pro Arg Gln His His Val Cys Val Gly Gly His Leu 500 505 510
- Gly Thr Pro Lys Lys Gly Cys Gly Tyr Leu Ser Gly Pro Gly Arg Ser 515 520 525
- Val Arg Leu Pro Ile Asp Ala Asp Arg Arg Ile Leu Val Gly Thr Ala 530 535 540

- Ala Met Ala Thr Lys Glu Ser Thr Thr Ser Pro Ser Val Lys Arg Met 545 550 555 555
- Leu Val Asp Thr Gln Gly Thr Asp Gln Trp Ile Ser Ala Gly Lys Ala 565 570 575
- Gln Gly Arg Met Pro Pro Ala Glu Ser Ala Arg Cys Arg His Pro Arg 580 585 590
- Asp Arg His Ser Ala Ser Val Arg Arg Cys Ser Thr Arg Trp Pro Val 595 600 605
- Thr Arg Arg Arg Ser Arg Ser Val Ala Trp Pro Arg Pro Pro Ser Pro 610 615 620
- Thr Cys Arg Arg Arg His Asp Leu Pro Ala Val Ala Ala Gly Ala 625 630 635 640
- Thr Ser Asn Trp Pro Ser Gly Lys Ala Thr Arg Pro Pro Thr Pro Pro 645 650 655
- Arg Trp Ala Ala Arg Gly Trp Pro Thr Leu Ala Gly Pro Leu Arg Ala 660 665 670
- Asp Ala Ala Cys Arg Ser Pro Val Ala Pro Thr Gly Phe Arg Pro 675 680 685
- Asp Pro Asp Ala Ile His Arg Cys Trp Pro Ala Gly Gln Ser Ala Ala 690 695 700
- Ala Ile Ala Ala Leu Val Ala Arg Tyr Pro Asp Ala Glu Thr Val Gln 705 710 715 720
- Leu His Pro Ala Asp Val Pro Phe Phe Val Thr Leu Cys Lys Thr Leu 725 730 735
- Gly Lys Pro Val Asn Phe Val Pro Ala Ile Asp Leu Val Val Arg Ala 740 745 750
- Gly Gly Ala Ala Thr Arg Cys Gly Arg Pro Thr Thr Pro Ala Thr Thr 755 760 765
- Pro Met Arg Cys Ala Ser Phe Arg Ala Arg Val Gly Ser Arg Ile Thr 770 780

Arg Met Asp Glu Pro Val Gly Glu Leu Leu Asp Ala Phe Glu Gln Ala 785 790 795 800

Ala Ile Asp Glu Val Leu Gly Ala Gly Val Glu Pro Lys Asp Val Ala 805 810 815

Ser Gly Arg Leu Gly Arg Ala Asp Val Ala Gly Pro Leu Ala Val Val 820 825 830

Leu Asp Ala Pro Asp Val Arg Trp Ala Gly Arg Thr Val Thr Asn Pro 835 840 845

Val His Arg Ile Arg Asp Pro Ala Glu Trp Gln Val His Asp Gly Pro 850 855 860

Glu Asn Pro Arg Ala Ala His Ser Ser Thr Gly Ala Arg Leu Gln Thr 865 870 875 880

His Gly Asp Asp Val Ala Leu Ser Val Ala Arg Leu Gly His Leu Gly 885 890 895

Arg His Pro Ile His Val Ala Gly Gln His Arg Arg Trp Arg His Pro $900 \hspace{1.5cm} 905 \hspace{1.5cm} 910$

Gly Asp Arg His Arg Gly Arg His His Ala Met Arg Thr Val Leu Arg 915 920 925

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Asp Gly His Phe Asp Gly Gly Leu Ala Pro Arg Ala Cys Cys Arg Pro 945 950 955 960

His Arg His Arg His Val Arg Cys Ala Leu Ala Pro Ser Leu Thr Asn 965 970 975

Val Pro Thr Arg Leu Val Gly Pro Cys Trp Pro Ala Val Phe Ala Ala 980 985 990

Ile Gly Ser Ala Val Thr Asp Thr Gly Glu Pro Val Val Glu Gly Leu 995 1000 1005

Leu Ser Leu Val His Leu Asp Thr Arg Pro Arg Val Val Gly Gln 1010 1015 1020

- Leu Pro Thr Val Pro Ala Gln Leu Thr Val Thr Gln Arg Leu Pro 1025 1030 1035
- Thr Gln Pro Ile Arg Thr Trp Ala Ala Ser Cys Arg Ser Arg Ser 1040 1045 1050
- Ser Phe Thr Ala Trp Arg Arg Asp Arg His Ser Arg Gly Ala Ile 1055 1060 1065
- Pro Val Ala Arg Cys Arg Glu Arg His Arg His Pro Ala Arg Arg 1085 1090 1095
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- Lys Met His Asp Ile Val Pro Arg Asp Glu Leu Gly Arg Ser Asn 1355 1360 1365
- Tyr Arg Leu Ser Ala Ile Arg Pro Ser Gln Ile Asp Leu Asp Asp 1370 1375 1380
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- Gln Thr Gly Ala His Leu Gln Arg Tyr Gly Pro Val Leu Ser Asp 1730 1735 1740
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Pro Glu Gly Arg Tyr Thr Phe Gly Ala Ser Cys Val Thr Thr Cys Pro 290 295 300

Tyr Asn Tyr Leu Ser Thr Glu Val Gly Ser Cys Thr Leu Val Cys Pro 305 310 315 320

Pro Asn Asn Glu Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu 325 330 335

Lys Cys Ser Lys Pro Cys Ala Arg Val Cys Tyr Gly Leu Gly Met Glu 340 345 350

His Leu Arg Gly Ala Arg Ala Ile Thr Ser Asp Asn Val Gln Glu Phe 355 360 365

Asp Gly Cys Lys Lys Ile Phe Gly Ser Leu Ala Phe Leu Pro Glu Ser 370 380

Phe Asp Gly Asp Pro Ser Ser Gly Ile Ala Pro Leu Arg Pro Glu Gln 385 390 395 400

Leu Gln Val Phe Glu Thr Leu Glu Glu Ile Thr Gly Tyr Leu Tyr Ile 405 410 415

Ser Ala Trp Pro Asp Ser Leu Arg Asp Leu Ser Val Phe Gln Asn Leu 420 425 430

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445

435 440

Leu Gln Gly Leu Gly Ile His Ser Leu Gly Leu Arg Ser Leu Arg Glu
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Val His Thr Val Pro Trp Asp Gln Leu Phe Arg Asn Pro His Gln Ala 485 490 495

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Gly Pro Thr Gln Cys Val Asn Cys Ser His Phe Leu Arg Gly Gln Glu 530 535 540

Cys Val Glu Glu Cys Arg Val Trp Lys Gly Leu Pro Arg Glu Tyr Val 545 550 555 560

Ser Asp Lys Arg Cys Leu Pro Cys His Pro Glu Cys Gln Pro Gln Asn 565 570 575

Ser Ser Glu Thr Cys Phe Gly Ser Glu Ala Asp Gln Cys Ala Ala Cys 580 585 590

Ala His Tyr Lys Asp Ser Ser Ser Cys Val Ala Arg Cys Pro Ser Gly 595 600

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Glu Gly Ile Cys Gln Pro Cys Pro Ile Asn Cys Thr His Ser Cys Val 625 630 635 640

Asp Leu Asp Glu Arg Gly Cys Pro Ala Glu Gln Arg Ala Ser Pro Val 645 650 655

Thr Phe Ile Ile Ala Thr Val Glu Gly Val Leu Leu Phe Leu Ile Leu 660 665 670

Val Val Val Gly Ile Leu Ile Lys Arg Arg Gln Lys Ile Arg

675

685

Lys Tyr Thr Met Arg Arg Leu Leu Gln Glu Thr Glu Leu Val Glu Pro $690 \hspace{1.5cm} 695 \hspace{1.5cm} 700 \hspace{1.5cm} .$

680

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Lys Glu Thr Glu Leu Arg Lys Val Lys Val Leu Gly Ser Gly Ala Phe
725 730 735

Gly Thr Val Tyr Lys Gly Ile Trp Ile Pro Asp Gly Glu Asn Val Lys
740 745 750

Ile Pro Val Ala Ile Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala 755 760 765

Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser 770 780

Pro Tyr Val Ser Arg Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln 785 790 795 800

Leu Val Thr Gln Leu Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg 805 810 815

Glu His Arg Gly Arg Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Val 820 825 830

Gln Ile Ala Lys Gly Met Ser Tyr Leu Glu Asp Val Arg Leu Val His 835 840 845

Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro Asn His Val 850 855 860

Lys Ile Thr Asp Phe Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr 865 870 875 880

Glu Tyr His Ala Asp Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu 885 890 895

Glu Ser Ile Leu Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser 900 905 910

Tyr Gly Val Thr Val Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr

915 920 925

Asp Gly Ile Pro Ala Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu 930 935 940

Arg Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met 945 950 955 960

Val Lys Cys Trp Met Ile Asp Ser Glu Cys Arg Pro Arg Phe Arg Glu 965 970 975

Leu Val Ser Glu Phe Ser Arg Met Ala Arg Asp Pro Gln Arg Phe Val 980 985 990

Phe Tyr Arg Ser Leu Leu Glu Asp Asp Asp Met Gly Asp Leu Val 1010 1015 1020

Asp Ala Glu Glu Tyr Leu Val Pro Gln Gln Gly Phe Phe Ser Pro 1025 1030 1035

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Ser Ser Ser Thr Arg Ser Gly Gly Gly Glu Leu Thr Leu Gly Leu 1055 1060 1065

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Val Thr Lys Gly Leu Gln Ser Leu Ser Pro His Asp Leu Ser Pro 1100 1105 1110

Leu Gln Arg Tyr Ser Glu Asp Pro Thr Leu Pro Leu Pro Pro Glu 1115 1120 1125

Thr Asp Gly Tyr Val Ala Pro Leu Ala Cys Ser Pro Gln Pro Glu 1130 1135 1140

Tyr Val Asn Gln Ser Glu Val Gln Pro Gln Pro Pro Leu Thr Pro

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Ser	Pro 1220		Phe	Asp	Asn	Leu 122		г Ту	r Tr	p As	p Gl: 12		sn S	Ser	Ser	
Glu	Gln 1235		Pro	Pro	Pro	Ser 124		n Ph	e Gl	u Gl		r P 45	ro T	ľhr	Ala	
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gcc Ala	gcg	ctc	tcg	ctc	tgg	ccg	acg	agt	gga	gaa	atc	tgc	ggg	cca	a ggc	153
	Ala	Leu	Ser	Leu 25	Trp	Pro	Thr	ser	G1у 30	Glu	Ile	Cys	Gly	Pro 35	o GLY	
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Ile	gac Asp gtg	atc Ile atc	cgc Arg 40	25 aac Asn	gac Asp tac	tat Tyr	cag Gln cac	cag Gln 45	30 ctg Leu	aag Lys ctc	cgc Arg	ctg Leu tcc	gag Glu 50 aag	35 aac Asr	tgc	201 249
Ile acg Thr	gac Asp gtg Val	atc Ile atc Ile 55	cgc Arg 40 gag Glu	25 aac Asn ggc Gly tac	gac Asp tac Tyr	tat Tyr ctc Leu	cag Gln cac His 60	cag Gln 45 atc Ile	30 ctg Leu ctg ctg ctc	aag Lys ctc Leu acg	cgc Arg atc Ile	ctg Leu tcc Ser 65	gag Glu 50 aag Lys acc	aac Asr gcc Ala	c tgc n Cys c gag	

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					atg Met											441
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Arg Lys Arg Arg Asp Val Met Gln Val Ala Asn Thr Thr Met Ser Ser 740 745 750

Arg Ser Arg Asn Thr Thr Ala Ala Asp Thr Tyr Asn Ile Thr Asp Pro 755 760 765

Glu Glu Leu Glu Thr Glu Tyr Pro Phe Phe Glu Ser Arg Val Asp Asn 770 775 780

Lys Glu Arg Thr Val Ile Ser Asn Leu Arg Pro Phe Thr Leu Tyr Arg 785 790 795 800

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Cys Ala	Val Let 35	ı Ser	Pro T		le G O	ln A	sp A	sp Pr	to Ly 45		r Hi	s Glįn	
Asn Gly 50	Ser Pro	Gln	_	eu T 5	hr G	ly T	hr A	ala Gl 60		n Va	l Pr	o Glu	
Ser Leu 65	Asp Ly	s Leu	His V 70	al T	hr S	er 1		Arg Pr 75	o Gl	n Ty	r Va	l Arg 80	

- Ile Lys Asn Trp Gly Ser Gly Glu Ile Leu His Asp Thr Leu His His 85 90 95
- Lys Ala Thr Ser Asp Phe Thr Cys Lys Ser Lys Ser Cys Leu Gly Ser 100 105 110
- Ile Met Asn Pro Lys Ser Leu Thr Arg Gly Pro Arg Asp Lys Pro Thr 115 120 125
- Pro Leu Glu Glu Leu Leu Pro His Ala Ile Glu Phe Ile Asn Gln Tyr 130 140
- Tyr Gly Ser Phe Lys Glu Ala Lys Ile Glu Glu His Leu Ala Arg Leu 145 150 155 160
- Glu Ala Val Thr Lys Glu Ile Glu Thr Thr Gly Thr Tyr Gln Leu Thr 165 170 175
- Leu Asp Glu Leu Ile Phe Ala Thr Lys Met Ala Trp Arg Asn Ala Pro 180 185 190
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- Arg Asn Cys Ser Thr Ala Gln Glu Met Phe Gln His Ile Cys Arg His $210 \\ \hspace{1.5cm} 215 \\ \hspace{1.5cm} 220 \\ \hspace{1.5cm}$
- Ile Leu Tyr Ala Thr Asn Asn Gly Asn Ile Arg Ser Ala Ile Thr Val 235 230 235
- Gln Leu Ile Arg Tyr Ala Gly Tyr Gln Met Pro Asp Gly Thr Ile Arg 260 265 270
- Gly Asp Ala Ala Thr Leu Glu Phe Thr Gln Leu Cys Ile Asp Leu Gly 275 280 285
- Trp Lys Pro Arg Tyr Gly Arg Phe Asp Val Leu Pro Leu Val Leu Gln 290 295 300
- Ala Asp Gly Gln Asp Pro Glu Val Phe Glu Ile Pro Pro Asp Leu Val 305 310 315 320

- Gly Leu Lys Trp Tyr Ala Leu Pro Ala Val Ala As
n Met Leu Leu Glu 340 345 350
- Val Gly Gly Leu Glu Phe Pro Ala Cys Pro Phe Asn Gly Trp Tyr Met 355 360 365
- Gly Thr Glu Ile Gly Val Arg Asp Phe Cys Asp Thr Gln Arg Tyr Asn 370 375 380
- Ile Leu Glu Glu Val Gly Arg Met Gly Leu Glu Thr His Thr Leu 385 390 395 400
- Ala Ser Leu Trp Lys Asp Arg Ala Val Thr Glu Ile Asn Val Ala Val 405 410 415
- Leu His Ser Phe Gln Lys Gln Asn Val Thr Ile Met Asp His His Thr 420 425 430
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- Val Lys Val Val Phe Phe Ala Ser Met Leu Met Arg Lys Val Met Ala 515 520 525
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- Glu Ala Leu Ala Arg Asp Leu Ala Thr Leu Phe Ser Tyr Ala Phe Asn 545 550 555 560

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Glu Gln Leu Leu Val Val Thr Ser Thr Phe Gly Asn Gly Asp Cys 585 590

Pro Ser Asn Gly Gln Thr Leu Lys Lys Ser Leu Phe Met Leu Arg Glu 595 600 605

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His Leu Gly Ala Ser Gln Leu Ala Pro Thr Gly Glu Gly Asp Glu Leu 645 650 655

Ser Gly Gln Glu Asp Ala Phe Arg Ser Trp Ala Val Gln Thr Phe Arg 660 665 670

Ala Ala Cys Glu Thr Phe Asp Val Arg Ser Lys His His Ile Gln Ile 675 680 685

Pro Lys Arg Phe Thr Ser Asn Ala Thr Trp Glu Pro Gln Gln Tyr Arg 690 695 700

Leu Ile Gln Ser Pro Glu Pro Leu Asp Leu Asn Arg Ala Leu Ser Ser 705 710 715 720

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Leu Gln Ser Glu Lys Ser Ser Arg Thr Thr Leu Leu Val Gln Leu Thr 740 745 750

Phe Glu Gly Ser Arg Gly Pro Ser Tyr Leu Pro Gly Glu His Leu Gly 755 760 765

Ile Phe Pro Gly Asn Gln Thr Ala Leu Val Gln Gly Ile Leu Glu Arg 770 775 780

Val Val Asp Cys Pro Thr Pro His Gln Thr Val Cys Leu Glu Val Leu 785 790 795 800

- Asp Glu Ser Gly Ser Tyr Trp Val Lys Asp Lys Arg Leu Pro Pro Cys 805 810 815
- Ser Leu Ser Gln Ala Leu Thr Tyr Phe Leu Asp Ile Thr Thr Pro Pro 820 825 830
- Thr Gln Leu Gln Leu His Lys Leu Ala Arg Phe Ala Thr Asp Glu Thr 835 840 845
- Asp Arg Gln Arg Leu Glu Ala Leu Cys Gln Pro Ser Glu Tyr Asn Asp 850 860
- Trp Lys Phe Ser Asn Asn Pro Thr Phe Leu Glu Val Leu Glu Glu Phe 865 870 875 880
- Pro Ser Leu His Val Pro Ala Ala Phe Leu Leu Ser Gln Leu Pro Ile 885 890 895
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										gag Glu					8052
										tac Tyr					8100
										act Thr 490					8148
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_		-								gga Gly					8244
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										tta Leu					8340
	 					_				gat Asp 570	_				8388
	 			_	-		_	-		tgg Trp			_		8436
_	_	_					-	_	_	ctc Leu					8484
										gaa Glu					8532
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					tac Tyr											8772
					tgc Cys											8820
		_	_		gtt Val		_	_		_			_	-	-	8868
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					aca Thr											8964
					atg Met						_					9012
					ggc Gly											9060
_	- 1				tgg Trp	-				_		_				9108
					acg Thr 820											9156
					atc Ile											9204
					ggg Gly											9252
					acg Thr											9300

cat His	aga Arg 880	cac His	agc Ser	acg Thr	ttc Phe	att Ile 885	cca Pro	ggg Gly	aca Thr	aat Asn	aaa Lys 890	Trp	gca Ala	a ggt a Gly	ctġ Leu	9348
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tgc Cys	tac Tyr	aca Thr	atg Met	aat Asn 915	cca Pro	aga Arg	aaa Lys	ctt Leu	ttt Phe 920	gac Asp	tac Tyr	tgt Cys	gat Ası	t atc p Ile 925	cct Pro	9444
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tgt Cys 975	gga Gly	ggc Gly	acc Thr	tta Leu	ata Ile 980	tcc Ser	cca Pro	gag Glu	tgg Trp	gto Val 985	Lei	g act 1 Thi	gc Al	t gct a Ala	cac His 990	9636
tgc Cys	ttg Leu	aag Lys	aag Lys	tcc Ser 995	tca Ser	agg Arg	cct Pro	tca Ser	tcc Ser 100	Тy	ıc aa r Ly	ag gt 7s Vá	c a al I	tc ct le Le 10	g ggt u Gly 05	9684
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	aag Lys		_	Ar	g cc g Pr	t gc o Al	c gt a Va	1 Il	са е Т 4 5			aaa (Lys '			cca Pro	9819
gct Ala	tgt Cys	ctg Leu	cca Pro 105	Se	c cc r Pr	a ga o As	с ta р Ту	r Me	g g t V 60	tc a al 1	acc (Thr)	gcc a Ala i	agg Arg	act Thr 1065	gaa Glu	9864
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tgc Cvs	+	cac	tat	. aa	a ta	t at	t tg	rt go	et c	gag (cat	ttg	gcc	aga	ggc	9999

act gac agt tgc cag ggt gac agt gga ggg cct ctg gtt tgc ttc Thr Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Phe 1115 1120 1125	10044
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Pro Ser Leu Glu Ala Pro Ser Glu Gln Ala Pro Thr Glu Gln Arg Pro 50 55 60	
Gly Val Gln Glu Cys Tyr His Gly Asn Gly Gln Ser Tyr Gln Gly Thr 65 75 80	
Tyr Phe Ile Thr Val Thr Gly Arg Thr Cys Gln Ala Trp Ser Ser Met 85 90 95	
Thr Pro His Ser His Ser Arg Thr Pro Ala Tyr Tyr Pro Asn Ala Gly 100 105 110	
Leu Ile Lys Asn Tyr Cys Arg Asn Pro Asp Pro Val Ala Ala Pro Trp 115 120 125	

- Cys Tyr Thr Thr Asp Pro Ser Val Arg Trp Glu Tyr Cys Asn Leu Thr 130 135 140
- Arg Cys Ser Asp Ala Glu Trp Thr Ala Phe Val Pro Pro Asn Val Ile 145 150 155 160
- Leu Ala Pro Ser Leu Glu Ala Phe Phe Glu Gln Ala Leu Thr Glu Glu 165 170 175
- Thr Pro Gly Val Gln Asp Cys Tyr Tyr His Tyr Gly Gln Ser Tyr Arg 180 185 190
- Gly Thr Tyr Ser Thr Thr Val Thr Gly Arg Thr Cys Gln Ala Trp Ser 195 200 205
- Ser Met Thr Pro His Gln His Ser Arg Thr Pro Glu Asn Tyr Pro Asn 210 215 220
- Ala Gly Leu Thr Arg Asn Tyr Cys Arg Asn Pro Asp Ala Glu Ile Arg 225 230 230 235
- Pro Trp Cys Tyr Thr Met Asp Pro Ser Val Arg Trp Glu Tyr Cys Asn 245 250 255
- Leu Thr Gln Cys Leu Val Thr Glu Ser Ser Val Leu Ala Thr Leu Thr 260 265 270
- Val Val Pro Asp Pro Ser Thr Glu Ala Ser Ser Glu Glu Ala Pro Thr 275 280 285
- Glu Gln Ser Pro Gly Val Gln Asp Cys Tyr His Gly Asp Gly Gln Ser 290 295 300
- Tyr Arg Gly Ser Phe Ser Thr Thr Val Thr Gly Arg Thr Cys Gln Ser 305 310 315 320
- Trp Ser Ser Met Thr Pro His Trp His Gln Arg Thr Thr Glu Tyr Tyr 325 330 335
- Pro Asn Gly Gly Leu Thr Arg Asn Tyr Cys Arg Asn Pro Asp Ala Glu 340 345 350
- Ile Ser Pro Trp Cys Tyr Thr Met Asp Pro Asn Val Arg Trp Glu Tyr 355 360 365

Cys Asn Leu Thr Gln Cys Pro Val Thr Glu Ser Ser Val Leu Ala Thr 370 375 380

Ser Thr Ala Val Ser Glu Gln Ala Pro Thr Glu Gln Ser Pro Thr Val 385 390 395 400

Gln Asp Cys Tyr His Gly Asp Gly Gln Ser Tyr Arg Gly Ser Phe Ser 405 410 415

Thr Thr Val Thr Gly Arg Thr Cys Gln Ser Trp Ser Ser Met Thr Pro 420 425 430

His Trp His Gln Arg Thr Thr Glu Tyr Tyr Pro Asn Gly Gly Leu Thr 435 440 445

Thr Met Asp Pro Ser Val Arg Trp Glu Tyr Cys Asn Leu Thr Gln Cys 465 470 475 480

Pro Val Met Glu Ser Thr Leu Leu Thr Thr Pro Thr Val Val Pro Val 485 490 495

Pro Ser Thr Glu Leu Pro Ser Glu Glu Ala Pro Thr Glu Asn Ser Thr 500 505 510

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Leu Ser Thr Thr Ile Thr Gly Arg Thr Cys Gln Ser Trp Ser Ser Met 530 540

Thr Pro His Trp His Arg Arg Ile Pro Leu Tyr Tyr Pro Asn Ala Gly 545 550 555 560

Leu Thr Arg Asn Tyr Cys Arg Asn Pro Asp Ala Glu Ile Arg Pro Trp 565 570 575

Cys Tyr Thr Met Asp Pro Ser Val Arg Trp Glu Tyr Cys Asn Leu Thr 580 595

Arg Cys Pro Val Thr Glu Ser Ser Val Leu Thr Thr Pro Thr Val Ala 595 600 605

Pro Val Pro Ser Thr Glu Ala Pro Ser Glu Gln Ala Pro Pro Glu Lys 610 615 620

Ser Pro Val Val Gln Asp Cys Tyr His Gly Asp Gly Arg Ser Tyr Arg 625 630 635 640

Gly Ile Ser Ser Thr Thr Val Thr Gly Arg Thr Cys Gln Ser Trp Ser 645 650 655

Ser Met Ile Pro His Trp His Gln Arg Thr Pro Glu Asn Tyr Pro Asn 660 665 670

Ala Gly Leu Thr Glu Asn Tyr Cys Arg Asn Pro Asp Ser Gly Lys Gln 675 680 685

Pro Trp Cys Tyr Thr Thr Asp Pro Cys Val Arg Trp Glu Tyr Cys Asn 690 695 700

Leu Thr Gln Cys Ser Glu Thr Glu Ser Gly Val Leu Glu Thr Pro Thr 705 710 715 720

Val Val Pro Val Pro Ser Met Glu Ala His Ser Glu Ala Ala Pro Thr $725 \hspace{1.5cm} 730 \hspace{1.5cm} 735$

Tyr Arg Gly Thr Phe Ser Thr Thr Val Thr Gly Arg Thr Cys Gln Ser 755 765

Trp Ser Ser Met Thr Pro His Arg His Gln Arg Thr Pro Glu Asn Tyr 770 775 780

Pro Asn Asp Gly Leu Thr Met Asn Tyr Cys Arg Asn Pro Asp Ala Asp 785 790 795 800

Thr Gly Pro Trp Cys Phe Thr Met Asp Pro Ser Ile Arg Trp Glu Tyr 805 810 815

Cys Asn Leu Thr Arg Cys Ser Asp Thr Glu Gly Thr Val Val Ala Pro $$ 820 $$ 825 $$ 830

- Cys Met Phe Gly Asn Gly Lys Gly Tyr Arg Gly Lys Lys Ala Thr Thr 850 855 860
- Val Thr Gly Thr Pro Cys Gln Glu Trp Ala Ala Gln Glu Pro His Arg 865 870 875 880
- His Ser Thr Phe Ile Pro Gly Thr Asn Lys Trp Ala Gly Leu Glu Lys 885 890 895
- Asn Tyr Cys Arg Asn Pro Asp Gly Asp Ile Asn Gly Pro Trp Cys Tyr 900 905 910
- Thr Met Asn Pro Arg Lys Leu Phe Asp Tyr Cys Asp Ile Pro Leu Cys 915 920 925
- Ala Ser Ser Ser Phe Asp Cys Gly Lys Pro Gln Val Glu Pro Lys Lys 930 935 940
- Cys Pro Gly Ser Ile Val Gly Gly Cys Val Ala His Pro His Ser Trp 945 950 955 960
- Pro Trp Gln Val Ser Leu Arg Thr Arg Phe Gly Lys His Phe Cys Gly 965 970 975
- Gly Thr Leu Ile Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu 980 985 990
- Lys Lys Ser Ser Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly Ala His 995 1000 1005
- Gln Glu Val Asn Leu Glu Ser His Val Gln Glu Ile Glu Val Ser 1010 1015 1020
- Arg Leu Phe Leu Glu Pro Thr Gln Ala Asp Ile Ala Leu Leu Lys 1025 1030 1035
- Leu Ser Arg Pro Ala Val Ile Thr Asp Lys Val Met Pro Ala Cys 1040 1045 1050
- Leu Pro Ser Pro Asp Tyr Met Val Thr Ala Arg Thr Glu Cys Tyr 1055 1060 1065
- Ile Thr Gly Trp Gly Glu Thr Gln Gly Thr Phe Gly Thr Gly Leu 1070 1075 1080

Leu	Lys 1085		Ala	Gln	Leu	Leu 109		1 11	e Gl	u As		u V 95	al C	Cys I	Asņ	
His	Tyr 1100		Туг	Ile	Cys	Ala 110		u Hi	s Le	u Al		g G 10	sly T	hr i	Asp	
Ser	Cys 1115		Gly	Asp	Ser	Gly 112		y Pr	o Le	u Va		s F 25	he G	Glu i	Lys	•
Asp	Lys 1130		Ile	Leu	Gln	Gly 113		l Th	ır Se	er Tr		y I 40	eu G	Sly	Cys	
Ala	Arg 1145		Asn	Lys	Pro	Gly 115		1 Ту	r Al	a Ar		ıl S .55	Ser A	Arg	Phe	
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acc Thr	cga Arg 45	ttc Phe	aag Lys	cac His	ctc Leu	cgg Arg 50	aag Lys	tac Tyr	aca Thr	tac Tyr	aac Asn 55	tat Tyr	gag Glu	gct Ala	gag a Glu	255
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ctg Leu	aag Lys	acc Thr	agc Ser 95	cag Gln	tgc Cys	atc Ile	ctg Leu	aaa Lys 100	gag Glu	gtg Val	tat Tyr	ggc Gly	ttc Phe 105	Asr	c cct n Pro	399
gaq	aac	aaa	acc	tta	ctq	aaq	aaa	acc	aaq	aac	tct	gag	gag	ttt	t gct	447

Glu	Gly	Lys 110	Ala	Leu	Leu	Lys	Lys 115	Thr	Lys	Asn	Ser	Glu 120	Glu	Phe	Ala	
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	cac His															687
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					gtg Val											1311
					ctg Leu											1359
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					gtc Val											1455
					gac Asp 465											1503
					gat Asp											1551
					caa Gln											1599
					tgt Cys											1647
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											gcc Ala					2127
		_						_	_		aaa Lys 695					2175
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-	ctg Leu 100	Ar			a gga r Gl		ú I			ag t In T	yr S		gtc Val	_	_	3132
acc Thr	tat Tyr 102	Gl	g ct u Le	c ca u Gl	g aga n Ar	a ga g Gl 10	u A	ac a .sp A	ga g rg A	cc t la I	eu V	tg al .030	gat Asp	acc Thr	ctg Leu	3177
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-	aca Thr 105	Ph	ć aa e Ly	a ta s Ty	t aa r As:	t cg n Ar 10	g G	ag a In S	gt a Ger M	itg a Met T	hr I	tg Leu 1060	tcc Ser	agt Ser	gaa Glu	3267
gtc	caa	at	t cc	g ga	t tt	t ga	t g	rtt g	ac c	etc g	ıga a	ıca	atc	ctc	aga	3312

Val	Gln 1065	Ile	Pro	Asp	Phe	Asp 1070	Val	Asp	Leu	Gly	Thr 1075	Ile	Leu	Arg	
gtt Val		gat Asp	gaa Glu	tct Ser	act Thr	gag	ggc Gly	aaa Lys	acg Thr	tct Ser	tac Tyr 1090	aga Arg	ctc Leu	acc Thr	3357
_	gac Asp 1095	att Ile	cag Gln	aac Asn	aag Lys	aaa Lys 1100	att Ile	act Thr	gag Glu	gtc Val	gcc Ala 1105	ctc Leu	atg Met		3402
	cta Leu 1110	agt Ser	tgt Cys	gac Asp	aca Thr	aag Lys 1115	gaa Glu	gaa Glu	aga Arg	aaa Lys	atc Ile 1120	aag Lys	ggt Gly		3447
	tcc Ser 1125	ata Ile	ccc Pro	cgt Arg	ttg Leu	caa Gln 1130	gca Ala	gaa Glu	gcc Ala	aga Arg	agt Ser 1135	gag Glu	atc Ile	ctc Leu	3492
_	cac His 1140	tgg Trp	tcg Ser	cct Pro	gcc Ala	aaa Lys 1145	ctg Leu	ctt Leu	ctc Leu	caa Gln	atg Met 1150	gac Asp	tca Ser	tct Ser	3537
	aca Thr 1155	gct Ala	tat Tyr	ggc Gly	tcc Ser	aca Thr 1160	gtt Val	tcc Ser	aag Lys	agg Arg	gtg Val 1165	gca Ala		cat His	3582
	gat Asp 1170										aca Thr 1180			aat Asn	3627
-	gat Asp 1185					act Thr 1190					gtg Val 1195				3672
						cat His 1205					aga Arg 1210				3717
		Val				gac Asp 1220		act Thr	ttc Phe	cgg Arg	cac His 1225	gtg Val		tcc Ser	3762
	tta Leu 1230	Ile	gtt Val	gca Ala	atg Met	agc Ser 1235					aag Lys 1240				3807
	ctt Leu 1245	Pro				act Thr 1250	ttg Leu	caa Gln	gac Asp	cac His	ctc Leu 1255	Asn	agc Ser	ctg Leu	3852
_	gag Glu 1260	Phe				aac Asn 1265	Met	gga Gly	ttg Leu	cca Pro	gac Asp 1270	Phe	cac His	atc Ile	3897
	gaa Glu 1275	Asn				aaa Lys 1280	Ser	gat Asp	ggc Gly	cgg Arg	gtc Val 1285	Lys	tat Tyr	acc Thr	3942
ttg	aac	aag	aac	agt	ttg	aaa	att	gag	att	cct	ttg	cct	ttt	ggt	3987

Leu	Asn 1290	Lys	Asn	Ser	Leu	Lys 1295	Ile	Glu	Ile	Pro	Leu 1300	Pro	Phe	Gly	
	aaa Lys 1305	tcc Ser	tcc Ser	aga Arg	gat Asp	cta Leu 1310	aag Lys	atg Met	tta Leu	gag Glu	act Thr 1315	gtt Val	agg Arg	aca Thr	4032
	gcc Ala 1320										ctg Leu 1330				4077
	ttc Phe 1335					ttt Phe 1340						tat Tyr	caa Gln	-	4122
caa Gln	gtg Val 1350	cct Pro	ctc Leu	ctg Leu	ggt Gly	gtt Val 1355	cta Leu	gac Asp	ctc Leu	tcc Ser	acg Thr 1360	aat Asn	gtc Val	tac Tyr	4167
	aac Asn 1365														4212
	aca Thr 1380					ctt Leu 1385						_	aag Lys		4257
gac Asp	tct Ser 1395	gtg Val	gtt Val	gac Asp	ctg Leu	ctt Leu 1400	Ser	tac Tyr	aat Asn	gtg Val	caa Gln 1405		tct Ser		4302
	aca Thr 1410					aag Lys 1415	Asn						tgt Cys		4347
	tct Ser 1425	Leu	cgc Arg	cac His	aaa Lys	ttt Phe 1430	Leu	gat Asp	tcg Ser	aat Asn	atc Ile 1435	aaa Lys	ttc Phe	agt Ser	4392
	gta Val 1440	Ğlu	aaa Lys	ctt Leu	gga Gly	aac Asn 1445	Asn	cca Pro	gtc Val	tca Ser	aaa Lys 1450	ggt Gly	tta Leu	cta Leu	4437
	ttc Phe 1455	gat Asp	gca Ala	tct Ser	agt Ser	tcc Ser 1460	tgg Trp	gga Gly	cca Pro	cag Gln	atg Met 1465	tct Ser	gct Ala		4482
	cat His 1470	Leu	gac Asp	tcc Ser	aaa Lys	aag Lys 1475	Lys	cag Gln	cat His	ttg Leu	ttt Phe 1480	gtc Val	aaa Lys	gaa Glu	4527
_	aag Lys 1485	Ile					Arg				ttc Phe 1495	Ţyr	gct Ala		4572
	aca Thr 1500	Tyr				tgt Cys 1505	Gln				aac Asn 1510			cgg Arg	4617
ctc	aat	gga	gag	tcc	aac	ctg	agg	ttt	aac	tcc	tcc	tac	ctc	caa	4662

I	eu	Asn 1515	Gly	Glu	Ser	Asn	Leu 1520	Arg	Phe	Asn	Ser	Ser 1525	Tyr	Leu	Gln	
_	-						gga Gly 1535									4 707
		acc Thr 1545					ctg Leu 1550									4752
		tcc Ser 1560					aac Asn 1565									4797
		aat. Asn 1575					aac Asn 1580						aag Lys		_	4842
	_	acc Thr 1590					aat Asn 1595									4887
-	•	gat Asp 1605					agg Arg 1610									4932
		aat Asn 1620					gag Glu 1625							ggc Gly		4977
-	•	aaa Lys 1635					gct Ala 1640									5022
		gat Asp 1650					agt Ser 1655									5067
		ctg Leu 1665					gag Glu 1670						ggc Gly			5112
-		gca Ala 1680					aca Thr 1685	Thr	aat Asn	ggc Gly	cgc Arg	ttc Phe 1690	agg Arg	gaa Glu	cac His	5157
							gat Asp 1700									5202
							cag Gln 1715						Val			5247
		aac Asn 1725	att Ile	ttc Phe	aac Asn	ttc Phe	aag Lys 1730	Val	agt Ser	caa Gln	gaa Glu	gga Gly 1735	Leu	aag Lys	ctc Leu	5292
1	tca	aat	gac	atg	atg	ggc	tca	tat	gct	gaa	atg	aaa	ttt	gac	cac	5337

Ser	Asn 1740	Asp	Met	Met	Gly	Ser 1745	Tyr	Ala	Glu	Met	Lys 1750	Phe	Asp	His	
aca Thr	aac Asn 1755					gca Ala 1760						ttc Phe			5382
	ctt Leu 1770			att Ile		agc Ser 1775					tat Tyr 1780	aag Lys	caa Gln	_	5427
_	aat Asn 1785					ccc Pro 1790	tat Tyr	tct Ser	ctg Leu	gta Val	act Thr 1795		tta Leu		5472
-	gac Asp 1800	ctg Leu	aaa Lys	tac Tyr	aat Asn	gct Ala 1805	ctg Leu	gat Asp	ctc Leu	acc Thr	aac Asn 1810		ggg Gly		5517
	cgg Arg 1815	cta Leu	gaa Glu	ccc Pro	ctg Leu	aag Lys 1820	ctg Leu	cat His	gtg Val	gct Ala	ggt Gly 1825	aac Asn			5562
	gcc Ala 1830	Tyr				gaa Glu 1835					tat Tyr 1840			tct Ser	5607
	gct Ala 1845	Āla				agc Ser 1850	tat Tyr	aaa Lys	gca Ala	gac Asp	act Thr 1855	gtt Val	gct Ala	aag Lys	5652
gtt Val	cag Gln 1860	Gly	gtg Val	gag Glu	ttt Phe	agc Ser 1865	His	cgg Arg	ctc Leu	aac Asn	aca Thr 1870	gac Asp	atc Ile	gct Ala	5697
	ctg Leu 1875	Ala	tca Ser	gcc Ala	att Ile	gac Asp 1880	atg Met	agc Ser	aca Thr	aac Asn	tat Tyr 1885	aat Asn	tca Ser	gac Asp	5742
	ctg Leu 1890	His				gtc Val 1895	Phe				atg Met 1900			ttt Phe	5787
	atg Met 1905	Thr	atc Ile	gat Asp	gca Ala	cat His 1910	aca Thr	aat Asn	ggc Gly	aat Asn	ggg Gly 1915			gct Ala	5832
ctc Leu	tgg Trp 1920	Gly	gaa Glu	cat His	act Thr	ggg Gly 1925	Gln	ctg Leu	tat Tyr	agc Ser	aaa Lys 1930	Phe		ttg Leu	5877
aaa Lys	gca Ala 1935	Glu	cct Pro	ctg Leu	gca Ala	ttt Phe 1940	Thr	ttc Phe	tct Ser	cat His	gat Asp 1945	Tyr	aaa Lys	ggc Gly	5922
	aca Thr 1950	Ser				gtg Val 1955	Ser	agg Arg	aaa Lys	agc Ser	atc Ile 1960	Ser	_	gct Ala	5967
ctt	gaa	cac	aaa	gtc	agt	gcc	ctg	ctt	act	cca	gct	gag	cag	aca	6012

Leu	Glu 1965	His	Lys	Val	Ser	Ala 1970	Leu	Leu	Thr	Pro	Ala 1975	Glu	Gln	Thr	
	acc Thr 1980					acc Thr 1985									6057
cag Gln		ttg Leu	gat Asp	gct Ala	tac Tyr	aac Asn 2000	act Thr	aaa Lys	gat Asp	aaa Lys	att Ile 2005	ggc Gly	gtg Val	gag Glu	6102
	act Thr 2010		_		_	gct Ala 2015						-			6147
	aaa Lys 2025												att Ile	-	6192
~ _	tta Leu 2040					gcc Ala 2045									6237
•	gtt Val 2055					tat Tyr 2060									6282
	aac Asn 2070					gag Glu 2075					tat Tyr 2080		gag Glu		6327
	cga Arg 2085												aga Arg		6372
_	aag Lys 2100					gat Asp 2105							aga Arg		6417
	ctg Leu 2115					cag Gln 2120		gct Ala	aat Asn	gat Asp	tat Tyr 2125	ctg Leu	aat Asn		6462
	aat Asn 2130	tgg Trp	gag Glu	aga Arg	caa Gln	gtt Val 2135	tca Ser	cat His	gcc Ala	aag Lys	gag Glu 2140	aaa Lys	ctg Leu	act Thr	6507
_	ctc Leu 2145					aga Arg 2150	_				gat Asp 2155		caa Gln		6552
-	tta Leu 2160	gat Asp	gat Asp	gcc Ala	aaa Lys	atc Ile 2165	aac Asn	ttt Phe	aat Asn	gaa Glu	aaa Lys 2170	Leu	tct Ser	caa Gln	6597
_	cag Gln 2175			_		caa Gln 2180					att Ile 2185				6642
tat	gat	tta	cat	gat	ttg	aaa	ata	gct	ạtt	gct	aat	att	att	gat	6687

Tyr	Asp 2190	Leu	His	Asp	Leu	Lys 2195	Ile	Ala	Ile	Ala	Asn 2200	Ile	Ile	Asp		
gaa Glu	atc Ile 2205					aaa Lys 2210						tat Tyr			1	6732
	gta Val 2220			gta Val		aca Thr 2225				cta Leu	cat His 2230	ttg Leu		_	i	6777
gaa Glu	aat Asn 2235	att Ile	gat Asp	ttt Phe	aac Asn	aaa Lys 2240	agt Ser	gga Gly	agt Ser	agt Ser	act Thr 2245	gca Ala	tcc Ser	tgg Trp		6822
	caa Gln 2250					aag Lys 2255										6867
	aaa Lys 2265					aag Lys 2270							gac Asp			6912
_	cac His 2280	cta Leu	gct Ala	gga Gly	aag Lys	tta Leu 2285	aaa Lys	caa Gln	cac	att Ile	gag Glu 2290	gct Ala	att Ile	gat Asp		6957
gtt Val	aga Arg 2295	gtg Val	ctt Leu	tta Lėu	gat Asp	caa Gln 2300	ttg Leu	gga Gly	act Thr	aca Thr	att Ile 2305	Ser	ttt Phe	gaa Glu		7002
-	ata Ile 2310	aat Asn	gat Asp	gtt Val	ctt Leu	gag Glu 2315	cat His	gtc Val	aaa Lys	cac His	ttt Phe 2320	gtt Val	ata Ile	aat Asn		7047
	att Ile 2325					gta Val 2330								aga Arg		7092
	aaa Lys 2340	gtc Val	cat His	gag Glu	tta Leu	atc Ile 2345	gag Glu	agg Arg	tat Tyr	gaa Glu	gta Val 2350	gac Asp		caa Gln		7137
	cag Gln 2355					aaa Lys 2360	tta Leu	gta Val	gag Glu	ttg Leu	gcc Ala 2365	cac His		tac Tyr		7182
aag Lys	ttg Leu 2370	Lys	gag Glu	act Thr	att Ile	cag Gln 2375	Lys	cta Leu	agc Ser	aat Asn	gtc Val 2380	Leu	caa Gln	caa Gln		7227
	aag Lys 2385	Ile	aaa Lys	gat Asp	tac Tyr	ttt Phe 2390	Glu	aaa Lys	ttg Leu	gtt Val	gga Gly 2395	Phe	att Ile	gat Asp		7272
	gct Ala 2400	Val	aag Lys	aag Lys	ctt Leu	aat Asn 2405	Glu	tta Leu	tct Ser	ttt Phe	aaa Lys 2410	Thr		att Ile		7317
gaa	gat	gtt	aac	aaa	ttc	ctt	gac	atg	ttg	ata	aag	aaa	tta	aag		7362

Glu	Asp 2415	Val	Asn	Lys	Phe	Leu 2420	Asp	Met	Leu	Ile	Lys 2425	Lys	Leu	Lys		
							Val				aat Asn 2440				740	7
_	gag Glu 2445										cag Gln 2455				7452	2
	cca Pro 2460										tta Leu 2470				749	7
	gcc Ala 2475										cag Gln 2485				754:	2
	acc Thr 2490										tta Leu 2500		tca Ser	-	758	7
	ttg Leu 2505										act Thr 2515		gaa Glu	-	763:	2
											cag Gln 2530		ctt Leu		767	7
											aca Thr 2545				772	2
						act Thr 2555					aac Asn 2560				776	7
	_					atc Ile 2570					aaa Lys 2575		atg Met		781	2
											atc Ile 2590				785	7
	ggg Gly 2595										cag Gln 2605				790	2
	gct Ala 2610	Thr					Asp				ccc Pro 2620	Leu			794	7
	agg Arg 2625	Ile				cag Gln 2630	Ile				gac Asp 2635	Leu			799	2
ata	aaa	atc	cça	tcc	agg	ttt	tcc	aca	cca	gaa	ttt	acc	atc	ctt	803	7

Ile	Lys 2640	Ile	Pro	Ser	Arg	Phe 2645	Ser	Thr	Pro	Glu	Phe 2650	Thr	Ile	Leu	
	acc Thr 2655										ttt Phe 2665				8082
	gta Val 2670	aag Lys	atc Ile	atc Ile	aga Arg	acc Thr 2675	att Ile	gac Asp	cag Gln	atg Met	ctg Leu 2680	aac Asn	agt Ser	gag Glu	8127
	cag Gln 2685										gat Asp 2695		aag Lys		8172
~ ~	gac Asp 2700										gac Asp 2710				8217
	gaa Glu 2715										act Thr 2725		aac Asn		8262
	gat Asp 2730	ttt Phe	caa Gln	gtt Val	cct Pro	gac Asp 2735	ctt Leu	cac His	ata Ile	cca Pro	gaa Glu 2740	ttc Phe	cag Gln		8307
	cac His 2745					att Ile 2750		_			ttt Phe 2755		aag Lys		8352
tac Tyr	agt Ser 2760	att Ile	ctg ·Leu	aaa Lys	atc Ile	caa Gln 2765	tct Ser	cct Pro	ctt Leu	ttc Phe	aca Thr 2770	tta Leu	gat Asp	gca Ala	8397
aat Asn	gct Ala 2775	gac Asp	ata Ile	ggg Gly	aat Asn	gga Gly 2780	acc Thr	acc Thr	tca Ser	gca Ala	aac Asn 2785	Glu		ggt Gly	8442
	gca Ala 2790					gcc Ala 2795					aaa Lys 2800	Leu	gaa Glu	gtt Val	8487
ctc Leu	aat Asn 2805	ttt Phe	gat Asp	ttt Phe	caa Gln	gca Ala 2810	aat Asn	gca Ala	caa Gln	ctc Leu	tca Ser 2815	Asn	cct Pro	aag Lys	8532
	aat Asn 2820	Pro				aag Lys 2825					ttc Phe 2830	Ser			8577
	ctg Leu 2835	Arg				ggg Gly 2840	Ser	_	-	_	ttt Phe 2845	Phe			8622
_	att Ile 2850	Ğlu				aac Asn 2855					tta Leu 2860	His		gaa Glu	8667
aaa	aat	aca	ctg	gag	ctt	agt	aat	gga	gtg	att	gtc	aag	ata	aac	8712

Lys	Asn 2865	Thr	Leu	Glu	Leu	Ser 2870	Asn	Gly	Val	Ile	Val 2875	Lys	Ile	Asn	
						agc Ser 2885							aaa Lys	_	8757
	atc Ile 2895					ttc Phe 2900						` -	cgc Arg		8802
, ,	atc Ile 2910	_		_	_	aaa Lys 2915	-				gca Ala 2920		act Thr		8847
						aaa Lys 2930		_	_		_			-	8892
	gga Gly 2940					caa Gln 2945							gga Gly		8937
	act Thr 2955					tcc Ser 2960							cac His		8982
	gta Val 2970					gtt Val 2975					tcc Ser 2980		aac Asn		9027
			-			tca Ser 2990		-	-		-		gtg Val		9072
	agt Ser 3000					aaa Lys 3005							gaa Glu		9117
	gca Ala 3015					agg Arg 3020							gga Gly	_	9162
	att Ile 3030					aat Asn 3035									9207
	gag Glu 3045		_	_		aca Thr 3050			_		aat Asn 3055	_	aaa Lys	-	9252
_	ttt Phe 3060					aca Thr 3065									9297
	gca Ala 3075	_		_	_	ccc Pro 3080	Ser	_	_		gca Ala 3085	-		caa Gln	9342
gta	agt	gct	agg	ttc	aat	cag	tat	aag	tac	aac	caa	aat	ttc	tct	9387

Val	Ser 3090	Ala	Arg	Phe	Asn	Gln 3095	Tyr	Lys	Tyr	Ásn	Gln 3100	Asn	Phe	Ser	
_	gga Gly 3105					att Ile 3110									9432
	gaa Glu 3120					ttc Phe 3125					tta Leu 3130		att Ile		9477
Ğlu						aca Thr 3140					cct Pro 3145		ctg Leu		9522
						aaa Lys 3155									9567
	aca Thr 3165					gat Asp 3170									9612
	aac Asn 3180					tçc Ser 3185					ttg Leu 3190		gtg Val		9657
						agc Ser 3200									9702
						gca Ala 3215							tcc Ser		9747
	gaa Glu 3225					ttt Phe 3230		_			gct Ala 3235	-	aaa Lys		9792
						acc Thr 3245					gga Gly 3250		act Thr		9837
	gtt Val 3255					gtg Val 3260					ata Ile 3265		atg Met	tcg Ser	9882
						cca Pro 3275					atg Met 3280				9927
	atc Ile 3285					gtc Val 3290					tac Tyr 3295		tta Leu		9972
_	cca Pro 3300					cca Pro 3305				_	cct Pro 3310	_			10017
aag	ctt	tct	ctt	сса	gat	ttc	aag	gaa	ttg	tgt	acc	ata	agc	cat	10062

Lys	Leu 3315	Ser	Leu	Pro	Asp	Phe 3320	Lys	Glu	Leu	Cys	Thr 3325	Ile	Ser	His	
	ttt Phe 3330					ggc Gly 3335									10107
	tca Ser 3345					ctg Leu 3350									10152
_	tca Ser 3360	_		-		cat His 3365							tct Ser	_	10197
						aaa Lys 3380						Arg	ttg Leu		10242
	aaa Lys 3390					tta Leu 3395							agc Ser		10287
	ttt Phe 3405	~ ~			_	cat His 3410		_			_		acc Thr		10332
	aat Asn 3420	_	_			gtg Val 3425	-						caa Gln		10377
	att Ile 3435					ttc Phe 3440							aat Asn		10422
-	tca Ser 3450					tct Ser 3455							tat Tyr		10467
	aat Asn 3465			_	_	tac Tyr 3470			_			_	_	_	10512
	aag Lys 3480					agc Ser 3485									10557
	tct Ser 3495					gtc Val 3500							cgg Arg		10602
	tca Ser 3510					agt Ser 3515									10647
_	agc Ser 3525					gtg Val 3530							aaa Lys		10692
gat	gat	atc	tgg	aac	ctt	gaa	gta	aaa	gaa	aat	ttt	gct	gga	gaa	10737

Asp	Asp 3540	Ile	Trp	Asn	Leu	Glu 3545	Val	Lys	Glu	Asn	Phe 3550	Ala	Gly	Glu	
gcc Ala	aca Thr 3555					tat Tyr 3560				_		-	acg Thr		10782
	cac His 3570					ggc Gly 3575						gga Gly	gaa Glu		10827
	agc Ser 3585		_		_	gaa Glu 3590						_	tca Ser	-	10872
	gtt Val 3600					agt Ser 3605							gat Asp		10917
	gac Asp 3615					gtg Val 3620							aag Lys		10962
_	aag Lys 3630		_			aat Asn 3635	_	_			cat His 3640		ggg Gly		11007
	cag Gln 3645	_	_	-		ctt Leu 3650					gaa Glu 3655	_	gca Ala		11052
	gac Asp 3660					tta Leu 3665									11097
	atc Ile 3675				_	tat Tyr 3680	_	_	_		tgg Trp 3685	_			11142
_	ctg Leu 3690					agc Ser 3695	Ile								11187
_	tca Ser 3705					tac Tyr 3710					aat Asn 3715			tca Ser	11232
	tcc Ser 3720					gtt Val 3725	Leu								11277
	ctg Leu 3735	Lys				cta Leu 3740	Asn					Met	cct Pro		11322
	cat His 3750	Val				gat Asp 3755	Leu					Cys			11367
gac	ttc	aga	gaa	ata	caa	atc	tat	aag	aag	ctg	aga	act	tca	tca	11412

Asp	Phe 3765	Arg	Glu	Ile	Gln	Ile 3770	Tyr	Lys	Lys	Leu	Arg 3775	Thr	Ser	Ser	
	gcc Ala 3780					aca Thr 3785					aaa Lys 3790		cct Pro		11457
gtt Val	_					tat Tyr ,3800				_	_		ttg Leu		11502
	ttt Phe 3810					gtg Val 3815							gtg Val		11547
_	ttc Phe 3825					agt Ser 3830						_	-	ttg Leu	11592
	cta Leu 3840					aac Asn 3845					ttt Phe 3850		ttg Leu		11637
	atc Ile 3855					cag Gln 3860							att Ile	_	11682
	tct Ser 3870					att Ile 3875					ttt Phe 3880		gca Ala		11727
	gca Ala 3885					gac Asp 3890					aat Asn 3895	-	act Thr		11772
_	gcc Ala 3900					aaa Lys 3905					gaa Glu 3910		gtc Val	ctg Leu	11817
_	tcc Ser 3915					acc Thr 3920					gaa Glu 3925		gaa Glu		11862
	gtt Val 3930					aaa Lys 3935					acg Thr 3940		gcc Ala		11907
	act Thr 3945	Lys					His				agt Ser 3955				11952
	gaa Glu 3960	Asp					Gly				tgg Trp 3970				11997
	cac His 3975	Leu				agc Ser 3980	Pro				gat Asp 3985	ctc Leu		_	12042
cgc	tac	cag	aaa	gac	aag	aaa	ggc	atc	tcc	acc	tca	gca	gcc	tcc	12087

Arg	Tyr 3990	Gln	Lys	Asp	Lỳs	Lys 3995	Gly	Ile	Ser	Thr	Ser 4000	Ala	Ala	Ser	
cca Pro						ggc Gly 4010						gat Asp			12132
	tct Ser 4020	aaa Lys	tgg Trp	aac Asn	ttc Phe	tac Tyr 4025	tac Tyr	agc Ser	cct Pro	cag Gln	tcc Ser 4030		cca Pro	_	12177
	aaa Lys 4035	ctc Leu	acc Thr	ata Ile	ttc Phe	aaa Lys 4040	act Thr	gag Glu	ttg Leu	agg Arg	gtc Val 4045	cgg Arg			12222
	gag Glu 4050					aaa Lys 4055								gct Ala	12267
	ggc Gly 4065	ttg Leu	cta Leu	acc Thr	tct Ser	ctg Leu 4070	aaa Lys	gac Asp	aac Asn	gtg Val	ccc Pro 4075	aag Lys	gcc Ala	aca Thr	12312
	gtc Val 4080					gtc Val 4085					tgg Trp 4090				12357
	ctc Leu 4095	Thr	ctg Leu	aga Arg	gaa Glu	gtg Val 4100	tct Ser	tca Ser	aag Lys	ctg Leu	aga Arg 4105	aga Arg	aat Asn	ctg Leu	12402
_	gac Asp 4110	_				gtt Val 4115									12447
-	gat Asp 4125	Ile	gac Asp	gag Glu	agg Arg	ttc Phe 4130	cag Gln	aaa Lys	gga Gly	gcc Ala	agt Ser 4135			act Thr	12492
	acc Thr 4140	Tyr	caa Gln	gag Glu	tgg Trp	aag Lys 4145	Asp	aag Lys	gcc Ala	cag Gln	aat Asn 4150	Leu		cag Gln	12537
gaa Glu	ctg Leu 4155	Leu	act Thr	cag Gln	gaa Glu	ggc Gly 4160	caa Gln	gcc Ala	agt Ser	ttc Phe	cag Gln 4165	бĺу		aag Lys	12582
_	aac Asn 4170	Val	ttt Phe	gat Asp	ggc Gly	ttg Leu 4175	Val	cga Arg	gtt Val	act Thr	caa Gln 4180	Ğlu		cat His	12627
	aaa Lys 4185	Val	aag Lys	cat His	ctg Leu	att Ile 4190	Asp	tca Ser	ctc Leu	att Ile	gat Asp 4195	Phe	ctg Leu	aac Asn	12672
	ccc Pro 4200	Arg		_		ccg Pro 4205	Gly				ata Ile 4210	Tyr		agg Arg	12717
gag	gaa	ctt	. tgc	act	atg	ttc	ata	agg	gag	gta	ggg	acg	gta	ctg	12762

Glu	Glu 4215	Leu	Cys	Thr	Met	Phe 4220	Ile	Arg	Glu	Val	Gly 4225	Thr	Val	Leu	
	cag Gln 4230	gta Val	tat Tyr	tcg Ser	aaa Lys	gtc Val 4235	cat His	aat Asn	ggt Gly	tca Ser	gaa Glu 4240		ctg Leu		12807
	tat Tyr 4245			gac Asp		gtg Val 4250					ttc Phe 4255	gag Glu	tta Leu		12852
	cat His 4260	aaa Lys	cta Leu	ata Ile	gat Asp	gta Val 4265	atc Ile	tcg Ser	atg Met	tat Tyr	agg Arg 4270	gaa Glu	_	ttg Leu	12897
	gat Asp 4275	tta Leu	tca Ser	aaa Lys	gaa Glu	gcc Ala 4280	caa Gln	gag Glu	gta Val	ttt Phe	aaa Lys 4285	gcc Ala	att Ile	cag Gln	12942
tct Ser	ctc Leu 4290					gtg Val 4295		cgt Arg	aat Asn	ctt Leu	cag Gln 4300	gac Asp		tta Leu	12987
	ttc Phe 4305	att Ile	ttc Phe	caa Gln	cta Leu	ata Ile 4310	gaa Glu	gat Asp	aac Asn	att Ile	aaa Lys 4315	cag Gln	ctg Leu		13032
gag Glu	atg Met 4320	aaa Lys	ttt Phe	act Thr	tat Tyr	ctt Leu 4325	att Ile	aat Asn	tat Tyr	atc Ile	caa Gln 4330	gat Asp		atc Ile	13077
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	gaa Glu 4350	Asn				aat Asn 4355	cțt Leu	cat His	aag Lys	ttc Phe	aat Asn 4360	Glu	ttt Phe	att Ile	13167
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	tac Tyr 4380	Ile	atg Met	gcc Ala	ctt Leu	cgt Arg 4385	gaa Glu	gaa Glu	tat Tyr	ttt Phe	gat Asp 4390	cca Pro	agt Ser	ata Ile	13257
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_	ctg Leu 4410	Ile	aag Lys	aac Asn	ctg Leu	tta Leu 4415	Val	gct Ala	ctt Leu	aag Lys	gac Asp 4420	Phe		tct Ser	13347
_	tat Tyr 4425	Ile				tct Ser 4430	Asn				caa Gln 4435	Leu		agt Ser	13392
caa	gtt	gag	caa	ttt	ctg	cac	aga	aat	att	cag	gaa	tat	ctt	. agc	13437

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Ile	ctt Leu 4455	acc Thr	gat Asp	cca Pro	gat Asp	gga Gly 4460	aaa Lys	ggg Gly	aaa Lys	gag Glu	aag Lys 4465	att Ile	gca Ala	gag Glu	13482
ctt Leu	tct Ser 4470	gcc Ala	act Thr	gct Ala	cag Gln	gaa Glu 4475	ata Ile	att Ile	aaa Lys	agc Ser	cag Gln 4480	gcc Ala	att Ile	gcg Ala	13527
	aag Lys 4485	aaa Lys	ata Ile	att Ile	tct Ser	gat Asp 4490	tac Tyr	cac His	cag Gln	cag Gln	ttt Phe 4495	aga Arg	tat Tyr	aaa Lys	13572
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	gct Ala 4515	Glu	tcc Ser	aaa Lys	aga Arg	ttg Leu 4520	Ile	gac Asp	ctg Leu	tcc Ser	att Ile 4525			tac Tyr	13662
	aca Thr 4530	Phe	ctg Leu	ata Ile	tac Tyr	atc Ile 4535	Thr	gag Glu	tta Leu	ctg Leu	aaa Lys 4540	Lys	ctg Leu	caa Gln	13707
	acc Thr 4545	Thr	gtc Val	atg Met	aac Asn	ccc Pro 4550	Tyr	atg Met	aag Lys	ctt Leu	gct Ala 4555	Pro	gga Gly	gaa Glu	13752
ctt act atc atc ctc taa ttttttaaaa gaaatcttca tttattcttc Leu Thr Ile Ile Leu 4560											13800				
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cat	acagt	ga g	ccag	cctt	g ca	gtagg	cag	taga	ctat	aa g	cagaa	gcac	ata	tgaactg	13920
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		Pro	Pro					Leu F						co Ala	
Leu	ı Leu	Leu	Leu 20	Leu	Leu	Ala (Ala <i>F</i> 25	Arg F	Ala G	Glu Gl	lu G1 30	u Me	et Leu	
Glı	ı Asn	Val 35	Ser	Leu	Val		Pro 1	Lys <i>I</i>	Asp <i>P</i>	Ala I	hr Ai	rg Ph	ne Ly	ys His	

Leu Arg Lys Tyr Thr Tyr Asn Tyr Glu Ala Glu Ser Ser Ser Gly Val50

Pro Gly Thr Ala Asp Ser Arg Ser Ala Thr Arg Ile Asn Cys Lys Val 65 70 75 80

Glu Leu Glu Val Pro Gln Leu Cys Ser Phe Ile Leu Lys Thr Ser Gln 85 90 95

Cys Ile Leu Lys Glu Val Tyr Gly Phe Asn Pro Glu Gly Lys Ala Leu 100 105 110

Leu Lys Lys Thr Lys Asn Ser Glu Glu Phe Ala Ala Met Ser Arg 115 120 125

Tyr Glu Leu Lys Leu Ala Ile Pro Glu Gly Lys Gln Val Phe Leu Tyr 130 135 140

Pro Glu Lys Asp Glu Pro Thr Tyr Ile Leu Asn Ile Lys Arg Gly Ile 145 150 150 160

Ile Ser Ala Leu Leu Val Pro Pro Glu Thr Glu Glu Ala Lys Gln Val 165 170 175

Leu Phe Leu Asp Thr Val Tyr Gly Asn Cys Ser Thr His Phe Thr Val 180 . 185 . 190

Lys Thr Arg Lys Gly Asn Val Ala Thr Glu Ile Ser Thr Glu Arg Asp 195 200 205

Leu Gly Gln Cys Asp Arg Phe Lys Pro Ile Arg Thr Gly Ile Ser Pro 210 215 220

Leu Ala Leu Ile Lys Gly Met Thr Arg Pro Leu Ser Thr Leu Ile Ser 225 230 235 240

Ser Ser Gln Ser Cys Gln Tyr Thr Leu Asp Ala Lys Arg Lys His Val 245 250 255

Ala Glu Ala Ile Cys Lys Glu Gln His Leu Phe Leu Pro Phe Ser Tyr 260 265 270

Lys Asn Lys Tyr Gly Met Val Ala Gln Val Thr Gln Thr Leu Lys Leu 275 280 285

- Glu Asp Thr Pro Lys Ile Asn Ser Arg Phe Phe Gly Glu Gly Thr Lys 290 295 300
- Lys Met Gly Leu Ala Phe Glu Ser Thr Lys Ser Thr Ser Pro Pro Lys 305 310 315 320
- Gln Ala Glu Ala Val Leu Lys Thr Val Gln Glu Leu Lys Lys Leu Thr $325 \hspace{1.5cm} 330 \hspace{1.5cm} 335$
- Ile Ser Glu Gln Asn Ile Gln Arg Ala Asn Leu Phe Asn Lys Leu Val340 345 350
- Thr Glu Leu Arg Gly Leu Ser Asp Glu Ala Val Thr Ser Leu Leu Pro 355 360 365
- Gln Leu Ile Glu Val Ser Ser Pro Ile Thr Leu Gln Ala Leu Val Gln 370 375 380
- Cys Gly Gln Pro Gln Cys Ser Thr His Ile Leu Gln Trp Leu Lys Arg 385 390 395 400
- Val His Ala Asn Pro Leu Leu Ile Asp Val Val Thr Tyr Leu Val Ala 405 410 415
- Leu Ile Pro Glu Pro Ser Ala Gln Gln Leu Arg Glu Ile Phe Asn Met 420 425 430
- Ala Arg Asp Gln Arg Ser Arg Ala Thr Leu Tyr Ala Leu Ser His Ala 435 440 445
- Val Asn Asn Tyr His Lys Thr Asn Pro Thr Gly Thr Gln Glu Leu Leu 450 455 460
- Asp Ile Ala Asn Tyr Leu Met Glu Gln Ile Gln Asp Asp Cys Thr Gly 465 470 475 480
- Asp Glu Asp Tyr Thr Tyr Leu Ile Leu Arg Val Ile Gly Asn Met Gly 485 490 495
- Gln Thr Met Glu Gln Leu Thr Pro Glu Leu Lys Ser Ser Ile Leu Lys 500 505 510
- Cys Val Gln Ser Thr Lys Pro Ser Leu Met Ile Gln Lys Ala Ala Ile 515 520 525

- Gln Ala Leu Arg Lys Met Glu Pro Lys Asp Lys Asp Gln Glu Val Leu 530 535 540
- Leu Gln Thr Phe Leu Asp Asp Ala Ser Pro Gly Asp Lys Arg Leu Ala 545 550 555 560
- Ala Tyr Leu Met Leu Met Arg Ser Pro Ser Gln Ala Asp Ile Asn Lys 565 570 575
- Val Ala Ser His Ile Ala Asn Ile Leu Asn Ser Glu Glu Leu Asp Ile 595 600 605
- Gln Asp Leu Lys Lys Leu Val Lys Glu Val Leu Lys Glu Ser Gln Leu 610 615 620
- Pro Thr Val Met Asp Phe Arg Lys Phe Ser Arg Asn Tyr Gln Leu Tyr 625 630 635 640
- Lys Ser Val Ser Ile Pro Ser Leu Asp Pro Ala Ser Ala Lys Ile Glu 645 650 655
- Gly Asn Leu Ile Phe Asp Pro Asn Asn Tyr Leu Pro Lys Glu Ser Met 660 665 670
- Leu Lys Thr Thr Leu Thr Ala Phe Gly Phe Ala Ser Ala Asp Leu Ile 675 680 685
- Glu Ile Gly Leu Glu Gly Lys Gly Phe Glu Pro Thr Leu Glu Ala Leu 690 695 700
- Phe Gly Lys Gln Gly Phe Phe Pro Asp Ser Val Asn Lys Ala Leu Tyr 705 710 715 720
- Trp Val Asn Gly Gln Val Pro Asp Gly Val Ser Lys Val Leu Val Asp 725 730 735
- His Phe Gly Tyr Thr Lys Asp Asp Lys His Glu Gln Asp Met Val Asn 740 745 750
- Gly Ile Met Leu Ser Val Glu Lys Leu Ile Lys Asp Leu Lys Ser Lys 755 760 765

- Glu Val Pro Glu Ala Arg Ala Tyr Leu Arg Ile Leu Gly Glu Glu Leu 770 775 780
- Gly Phe Ala Ser Leu His Asp Leu Gln Leu Leu Gly Lys Leu Leu Room 785 790 795 800
- Met Gly Ala Arg Thr Leu Gln Gly Ile Pro Gln Met Ile Gly Glu Val 805 810 815
- Ile Arg Lys Gly Ser Lys Asn Asp Phe Phe Leu His Tyr Ile Phe Met 820 825 830
- Glu Asn Ala Phe Glu Leu Pro Thr Gly Ala Gly Leu Gln Leu Gln Ile 835 840 845
- Ser Ser Ser Gly Val Ile Ala Pro Gly Ala Lys Ala Gly Val Lys Leu 850 860
- Glu Val Ala Asn Met Gln Ala Glu Leu Val Ala Lys Pro Ser Val Ser 865 870 875 880
- Val Glu Phe Val Thr Asn Met Gly Ile Ile Ile Pro Asp Phe Ala Arg 885 890 895
- Ser Gly Val Gln Met Asn Thr Asn Phe Phe His Glu Ser Gly Leu Glu 900 905 910
- Ala His Val Ala Leu Lys Pro Gly Lys Leu Lys Phe Ile Ile Pro Ser 915 920 925
- Pro Lys Arg Pro Val Lys Leu Leu Ser Gly Gly Asn Thr Leu His Leu 930 935 940
- Val Ser Thr Thr Lys Thr Glu Val Ile Pro Pro Leu Ile Glu Asn Arg 945 950 955 960
- Gln Ser Trp Ser Val Cys Lys Gln Val Phe Pro Gly Leu Asn Tyr Cys 965 970 975
- Thr Ser Gly Ala Tyr Ser Asn Ala Ser Ser Thr Asp Ser Ala Ser Tyr 980 985 990
- Tyr Pro Leu Thr Gly Asp Thr Arg Leu Glu Leu Glu Leu Arg Pro Thr 995 1000 1005

- Arg Glu Asp Arg Ala Leu Val Asp Thr Leu Lys Phe Val Thr Gln 1025 1030 1035
- Ala Glu Gly Ala Lys Gln Thr Glu Ala Thr Met Thr Phe Lys Tyr 1040 1045 1050
- Asn Arg Gln Ser Met Thr Leu Ser Ser Glu Val Gln Ile Pro Asp 1055 1060 1065
- Phe Asp Val Asp Leu Gly Thr Ile Leu Arg Val Asn Asp Glu Ser 1070 1075 1080
- Thr Glu Gly Lys Thr Ser Tyr Arg Leu Thr Leu Asp Ile Gln Asn 1085 1090 1095
- Lys Lys Ile Thr Glu Val Ala Leu Met Gly His Leu Ser Cys Asp 1100 1105 1110
- Thr Lys Glu Glu Arg Lys Ile Lys Gly Val Ile Ser Ile Pro Arg 1115 1120 1125
- Leu Gln Ala Glu Ala Arg Ser Glu Ile Leu Ala His Trp Ser Pro 1130 1135 1140
- Ala Lys Leu Leu Gln Met Asp Ser Ser Ala Thr Ala Tyr Gly 1145 1150 1155
- Ser Thr Val Ser Lys Arg Val Ala Trp His Tyr Asp Glu Glu Lys 1160 1165 1170
- Ile Glu Phe Glu Trp Asn Thr Gly Thr Asn Val Asp Thr Lys Lys 1175 1180 1185
- Met Thr Ser Asn Phe Pro Val Asp Leu Ser Asp Tyr Pro Lys Ser 1190 1195 1200
- Leu His Met Tyr Ala Asn Arg Leu Leu Asp His Arg Val Pro Gln 1205 1210 1215
- Thr Asp Met Thr Phe Arg His Val Gly Ser Lys Leu Ile Val Ala 1220 1225 1230

- Met Ser Ser Trp Leu Gln Lys Ala Ser Gly Ser Leu Pro Tyr Thr 1235 1240 1245
- Gln Thr Leu Gln Asp His Leu Asn Ser Leu Lys Glu Phe Asn Leu 1250 1260
- Gln Asn Met Gly Leu Pro Asp Phe His Ile Pro Glu Asn Leu Phe 1265 1270 1275
- Leu Lys Ser Asp Gly Arg Val Lys Tyr Thr Leu Asn Lys Asn Ser 1280 1285 1290
- Leu Lys Ile Glu Ile Pro Leu Pro Phe Gly Gly Lys Ser Ser Arg 1295 1300 1305
- Asp Leu Lys Met Leu Glu Thr Val Arg Thr Pro Ala Leu His Phe 1310 1315 1320
- Lys Ser Val Gly Phe His Leu Pro Ser Arg Glu Phe Gln Val Pro 1325 1330 1335

- Trp Ser Ala Ser Tyr Ser Gly Gly Asn Thr Ser Thr Asp His Phe 1370 1375 1380
- Ser Leu Arg Ala Arg Tyr His Met Lys Ala Asp Ser Val Val Asp 1385 1390 1395
- Leu Leu Ser Tyr Asn Val Gln Gly Ser Gly Glu Thr Thr Tyr Asp 1400 1405 1410
- His Lys Asn Thr Phe Thr Leu Ser Cys Asp Gly Ser Leu Arg His 1415 1420 1425
- Lys Phe Leu Asp Ser Asn Ile Lys Phe Ser His Val Glu Lys Leu 1430 1435 1440
- Gly Asn Asn Pro Val Ser Lys Gly Leu Leu Ile Phe Asp Ala Ser 1445 1450 1455

- Lys Lys Gln His Leu Phe Val Lys Glu Val Lys Ile Asp Gly 1475 1480 1485
- Gln Phe Arg Val Ser Ser Phe Tyr Ala Lys Gly Thr Tyr Gly Leu 1490 1495 1500
- Ser Cys Gln Arg Asp Pro Asn Thr Gly Arg Leu Asn Gly Glu Ser 1505 1510 1515
- Asn Leu Arg Phe Asn Ser Ser Tyr Leu Gln Gly Thr Asn Gln Ile 1520 1530
- Asp Leu Gln Ser Gly Ile Ile Lys Asn Thr Ala Ser Leu Lys Tyr $1550 \\ \hspace*{1.5cm} 1555 \\ \hspace*{1.5cm} 1560 \\ \hspace*{1.5cm}$
- Glu Asn Tyr Glu Leu Thr Leu Lys Ser Asp Thr Asn Gly Lys Tyr 1565 1570 1575
- Lys Asn Phe Ala Thr Ser Asn Lys Met Asp Met Thr Phe Ser Lys 1580 1585 1590
- Gln Asn Ala Leu Leu Arg Ser Glu Tyr Gln Ala Asp Tyr Glu Ser 1595 1600 1605
- Leu Arg Phe Phe Ser Leu Leu Ser Gly Ser Leu Åsn Ser His Gly 1610 1620
- Leu Glu Leu Asn Ala Asp Ile Leu Gly Thr Asp Lys Ile Asn Ser 1625 1630 1635
- Gly Ala His Lys Ala Thr Leu Arg Ile Gly Gln Asp Gly Ile Ser 1640 1645 1650
- Thr Ser Ala Thr Thr Asn Leu Lys Cys Ser Leu Leu Val Leu Glu 1655 1660 1665
- Asn Glu Leu Asn Ala Glu Leu Gly Leu Ser Gly Ala Ser Met Lys 1670 1680

- Leu Thr $\,$ Thr Asn Gly Arg Phe $\,$ Arg Glu His Asn Ala $\,$ Lys Phe Ser $\,$ 1685 $\,$ 1695
- Leu Asp Gly Lys Ala Ala Leu Thr Glu Leu Ser Leu Gly Ser Ala 1700 1705 1710
- Tyr Gln Ala Met Ile Leu Gly Val Asp Ser Lys Asn Ile Phe Asn 1715 1720 1725
- Phe Lys Val Ser Gln Glu Gly Leu Lys Leu Ser Asn Asp Met Met 1730 1740
- Gly Ser Tyr Ala Glu Met Lys Phe Asp His Thr Asn Ser Leu Asn 1745 . 1750 . 1755
- Ile Ala Gly Leu Ser Leu Asp Phe Ser Ser Lys Leu Asp Asn Ile 1760 1765 1770
- Tyr Ser Ser Asp Lys Phe Tyr Lys Gln Thr Val Asn Leu Gln Leu 1775 1780 1785
- Gln Pro Tyr Ser Leu Val Thr Thr Leu As
n Ser Asp Leu Lys Tyr 1790 1795 1800
- Asn Ala Leu Asp Leu Thr Asn Asn Gly Lys Leu Arg Leu Glu Pro 1805 1810 1815
- Leu Lys Leu His Val Ala Gly Asn Leu Lys Gly Ala Tyr Gln Asn 1820 1830
- Ala Ser Tyr Lys Ala Asp Thr Val Ala Lys Val Gln Gly Val Glu 1850 1860
- Phe Ser His Arg Leu Asn Thr Asp Ile Ala Gly Leu Ala Ser Ala 1865 1870 1875
- Ile Asp Met Ser Thr Asn Tyr Asn Ser Asp Ser Leu His Phe Ser 1880 1885 1890

- Ala His Thr Asn Gly Asn Gly Lys Leu Ala Leu Trp Gly Glu His 1910 1915 1920
- Thr Gly Gln Leu Tyr Ser Lys Phe Leu Leu Lys Ala Glu Pro Leu 1925 1930 1935
- Ala Phe Thr Phe Ser His Asp Tyr Lys Gly Ser Thr Ser His His 1940 1945 1950
- Leu Val Ser Arg Lys Ser Ile Ser Ala Ala Leu Glu His Lys Val 1955 1960 1965
- Ser Ala Leu Leu Thr Pro Ala Glu Gln Thr Gly Thr Trp Lys Leu 1970 1975 1980
- Lys Thr GÍn Phe Asn Asn Glu Tyr Ser Gln Asp Leu Asp Ala 1985 1990 1995
- Tyr Asn Thr Lys Asp Lys Ile Gly Val Glu Leu Thr Gly Arg Thr 2000 2005 2010
- Leu Leu Ser Glu Pro Ile Asn Ile Ile Asp Ala Leu Glu Met Arg 2030 2035 2040
- Asp Ala Val Glu Lys Pro Gln Glu Phe Thr Ile Val Ala Phe Val 2045 2050 2055
- Lys Tyr Asp Lys Asn Gln Asp Val His Ser Ile Asn Leu Pro Phe 2060 2065 2070
- Phe Glu Thr Leu Gln Glu Tyr Phe Glu Arg Asn Arg Gln Thr Ile 2075 2080 2085
- Ile Val Val Leu Glu Asn Val Gln Arg Asn Leu Lys His Ile Asn 2090 2095 2100
- Ile Asp Gln Phe Val Arg Lys Tyr Arg Ala Ala Leu Gly Lys Leu 2105 2110 2115
- Pro Gln Gln Ala Asn Asp Tyr Leu Asn Ser Phe Asn Trp Glu Arg 2120 2125 2130

- Gln Val Ser His Ala Lys Glu Lys Leu Thr Ala Leu Thr Lys Lys 2135 2140 2145
- Lys Ile Asn Phe Asn Glu Lys Leu Ser Gln Leu Gln Thr Tyr Met 2165 2170 2175
- Ile Gln Phe Asp Gln Tyr Ile Lys Asp Ser Tyr Asp Leu His Asp 2180 2185 2190
- Leu Lys Ser Leu Asp Glu His Tyr His Ile Arg Val Asn Leu Val 2210 2215 2220
- Lys Thr Ile His Asp Leu His Leu Phe Ile Glu Asn Ile Asp Phe 2225 2230 2235
- Asn Lys Ser Gly Ser Ser Thr Ala Ser Trp Ile Gln Asn Val Asp 2240 2245 2250
- Thr Lys Tyr Gln Ile Arg Ile Gln Ile Gln Glu Lys Leu Gln Gln 2255 2260 2265
- Leu Lys Arg His Ile Gln Asn Ile Asp Ile Gln His Leu Ala Gly 2270 2275 2280
- Asp Gln Leu Gly Thr Thr Ile Ser Phe Glu Arg Ile Asn Asp Val 2300 2310
- Leu Glu His Val Lys His Phe Val Ile Asn Leu Ile Gly Asp Phe 2315 2320 2325
- Glu Val Ala Glu Lys Ile Asn Ala Phe Arg Ala Lys Val His Glu 2330 2335 2340

- Asp Lys Leu Val Glu Leu Ala His Gln Tyr Lys Leu Lys Glu Thr 2360 2365 2370
- Ile Gln Lys Leu Ser Asn Val Leu Gln Gln Val Lys Ile Lys Asp 2375 2380 2385
- Tyr Phe Glu Lys Leu Val Gly Phe Ile Asp Asp Ala Val Lys Lys 2390 2395 2400
- Leu Asn Glu Leu Ser Phe Lys Thr Phe Ile Glu Asp Val Asn Lys 2405 2410 2415
- Phe Leu Asp Met Leu Ile Lys Lys Leu Lys Ser Phe Asp Tyr His 2420 2425 2430
- Gln Phe Val Asp Glu Thr Asn Asp Lys Ile Arg Glu Val Thr Gln 2435 2440 2445
- Arg Leu Asn Gly Glu Ile Gln Ala Leu Glu Leu Pro Gln Lys Ala 2450 2460
- Glu Ala Leu Lys Leu Phe Leu Glu Glu Thr Lys Ala Thr Val Ala 2465 2470 2475
- Val Tyr Leu Glu Ser Leu Gln Asp Thr Lys Ile Thr Leu Ile Ile 2480 2485 2490
- Asn Trp Leu Gln Glu Ala Leu Ser Ser Ala Ser Leu Ala His Met 2495 2500 2505
- Lys Ala Lys Phe Arg Glu Thr Leu Glu Asp Thr Arg Asp Arg Met 2510 2515 2520
- Tyr Gln Met Asp Ile Gln Gln Glu Leu Gln Arg Tyr Leu Ser Leu 2525 2530 2535
- Val Gly Gln Val Tyr Ser Thr Leu Val Thr Tyr Ile Ser Asp Trp 2540 2545 2550
- Trp Thr Leu Ala Ala Lys Asn Leu Thr Asp Phe Ala Glu Gln Tyr 2555 2560 2565
- Ser Ile Gln Asp Trp Ala Lys Arg Met Lys Ala Leu Val Glu Gln 2570 2580

- Gly Phe Thr Val Pro Glu Ile Lys Thr Ile Leu Gly Thr Met Pro 2585 2590 2595
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- Thr Pro Asp Phe Ile Val Pro Leu Thr Asp Leu Arg Ile Pro Ser 2615 2620 2625
- Arg Phe Ser Thr Pro Glu Phe Thr Ile Leu Asn Thr Phe His Ile 2645 2650 2655
- Pro Ser Phe Thr Ile Asp Phe Val Glu Met Lys Val Lys Ile Ile 2660 2665 2670
- Arg Thr Ile Asp Gln Met Leu Asn Ser Glu Leu Gln Trp Pro Val 2675 2680 2685
- Ala Arg Ile Thr Leu Pro Asp Phe Arg Leu Pro Glu Ile Ala Ile 2705 2710 2715
- Pro Glu $\,$ Phe Ile Ile Pro Thr $\,$ Leu Asn Leu Asn Asp $\,$ Phe Gln Val $\,$ 2720 $\,$ $\,$ 2730 $\,$
- Pro Asp Leu His Ile Pro Glu Phe Gln Leu Pro His Ile Ser His 2735 2740 2745
- Thr Ile Glu Val Pro Thr Phe Gly Lys Leu Tyr Ser Ile Leu Lys 2750 2755 2760
- Ile Gln Ser Pro Leu Phe Thr Leu Asp Ala Asn Ala Asp Ile Gly 2765 2770 2775
- Asn Gly Thr Thr Ser Ala Asn Glu Ala Gly Ile Ala Ala Ser Ile 2780 2785 2790
- Thr Ala Lys Gly Glu Ser Lys Leu Glu Val Leu Asn Phe Asp Phe 2795 2800 2805

- Gln Ala Asn Ala Gln Leu Ser Asn Pro Lys Ile Asn Pro Leu Ala 2810 2815 2820
- Leu Lys Glu Ser Val Lys Phe Ser Ser Lys Tyr Leu Arg Thr Glu 2825 2830 2835
- His Gly Ser Glu Met Leu Phe Phe Gly Asn Ala Ile Glu Gly Lys 2840 2845 2850
- Ser Asn Thr Val Ala Ser Leu His Thr Glu Lys Asn Thr Leu Glu 2855 2860 2865
- Leu Ser Asn Gly Val Ile Val Lys Ile Asn Asn Gln Leu Thr Leu 2870 2875 2880
- Asp Ser Asn Thr Lys Tyr Phe His Lys Leu Asn Ile Pro Lys Leu 2885 2890 2895
- Leu Lys Ala Gly His Ile Ala Trp Thr Ser Ser Gly Lys Gly Ser 2915 2920 2925
- Trp Lys Trp Ala Cys Pro Arg Phe Ser Asp Glu Gly Thr His Glu 2930 2935 2940
- Ser Gln Ile Ser Phe Thr Ile Glu Gly Pro Leu Thr Ser Phe Gly 2945 2950 2955
- Leu Ser Asn Lys Ile Asn Ser Lys His Leu Arg Val Asn Gln Asn 2960 2965 2970
- Leu Val Tyr Glu Ser Gly Ser Leu Asn Phe Ser Lys Leu Glu Ile 2975 2980 2985
- Gln Ser Gln Val Asp Ser Gln His Val Gly His Ser Val Leu Thr 2990 2995 3000
- Ala Lys Gly Met Ala Leu Phe Gly Glu Gly Lys Ala Glu Phe Thr 3005 3010 3015
- Gly Arg His Asp Ala His Leu Asn Gly Lys Val Ile Gly Thr Leu 3020 3030

- Lys Asn Ser Leu Phe Phe Ser Ala Gln Pro Phe Glu Ile Thr Ala 3035 3040 3040
- Ser Thr Asn Asn Glu Gly Asn Leu Lys Val Arg Phe Pro Leu Arg 3050 3060
- Ser Pro Ser Ala Gln Gln Ala Ser Trp Gln Val Ser Ala Arg Phe 3080 3085 3090
- Asn Gln Tyr Lys Tyr Asn Gln Asn Phe Ser Ala Gly Asn Asn Glu 3095 3100 3105
- Asn Ile Met Glu Ala His Val Gly Ile Asn Gly Glu Ala Asn Leu 3110 3115 3120
- Asp Phe Leu Asn Ile Pro Leu Thr Ile Pro Glu Met Arg Leu Pro 3125 3130 3135
- Tyr Thr $\$ Ile $\$ Ile Thr $\$ Thr $\$ Pro $\$ Pro $\$ Leu $\$ Lys $\$ Asp $\$ Phe $\$ Ser $\$ Leu $\$ Trp $\$ 3140 $\$ 3150
- Glu Lys Thr Gly Leu Lys Glu Phe Leu Lys Thr Thr Lys Gln Ser 3155 3160 3165
- Phe Asp Leu Ser Val Lys Ala Gln Tyr Lys Lys Asn Lys His Arg 3170 3175 3180
- His Ser Ile Thr Asn Pro Leu Ala Val Leu Cys Glu Phe Ile Ser 3185 3190 3195
- Gln Ser Ile Lys Ser Phe Asp Arg His Phe Glu Lys Asn Arg Asn 3200 3205 3210
- Asn Ala Leu Asp Phe Val Thr Lys Ser Tyr Asn Glu Thr Lys Ile 3215 3220 3225
- Lys Phe Asp Lys Tyr Lys Ala Glu Lys Ser His Asp Glu Leu Pro 3230 3240
- Arg Thr Phe Gln Ile Pro Gly Tyr Thr Val Pro Val Val Asn Val 3245 3250 3255

- Glu Val Ser Pro Phe Thr Ile Glu Met Ser Ala Phe Gly Tyr Val 3260 3265 3270
- Phe Pro Lys Ala Val Ser Met Pro Ser Phe Ser Ile Leu Gly Ser 3275 3280 3285
- Asp Val Arg Val Pro Ser Tyr Thr Leu Ile Leu Pro Ser Leu Glu 3290 3295 3300
- Leu Pro Val Leu His Val Pro Arg Asn Leu Lys Leu Ser Leu Pro 3305 3310 3315
- Asp Phe Lys Glu Leu Cys Thr Ile Ser His Ile Phe Ile Pro Ala 3320 3330
- Thr Leu Asn Thr Asn Ala Glu Leu Phe Asn Gln Ser Asp Ile Val3350 3355 3360
- Ala His Leu Leu Ser Ser Ser Ser Val Ile Asp Ala Leu Gln 3365 3370 3375
- Tyr Lys Leu Glu Gly Thr Thr Arg Leu Thr Arg Lys Arg Gly Leu 3380 3380 3390
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- Ser Val Ala Thr Thr Thr Lys Ala Gln Ile Pro Ile Leu Arg Met 3425 3430 3435
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- Ala Ser Glu Ala Asn Thr Tyr Leu Asn Ser Lys Ser Thr Arg Ser 3515 3520 3525
- Ser Val Lys Leu Gln Gly Thr Ser Lys Ile Asp Asp Ile Trp Asn 3530 3535 3540
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- Thr Ser Ile Gly Arg Arg Gln His Leu Arg Val Ser Thr Ala Phe 3695 3700 3705

- Val Tyr Thr Lys Asn Pro Asn Gly Tyr Ser Phe Ser Ile Pro Val 3710 3715 3720
- Lys Val Leu Ala Asp Lys Phe Ile Ile Pro Gly Leu Lys Leu Asn 3725 3730 3735
- Asp Leu Asn Ser Val Leu Val Met Pro Thr Phe His Val Pro Phe 3740 3745 3750
- Thr Asp Leu Gln Val Pro Ser Cys Lys Leu Asp Phe Arg Glu Ile 3755 3760 3765
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Ser Ser P 3		Ala G	lu Leu	Gln 40	Cys	Cys	Ala	Gly	Trp 45	Arg	Gln	Lys	
Asp Gln G 50	lu Cys	Thr I	le Pro 55	Ile	Cys	Glu	G1y	Pro 60	Asp	Ala	Cys	Gln	
Lys Asp G	lu Val		al Lys	Pro	Gly	Leu	Cys 75	Arg	Cys	Lys	Pro	Gly 80	

Pro Asp Cys Arg Glu Ser Cys Pro Cys His Pro His Gly Gln Cys Glu

Phe Phe Gly Ala His Cys Ser Ser Arg Cys Pro Gly Gln Tyr Trp Gly

100 105 110

Pro Ala Thr Gly Ala Cys Gln Cys Gln Ala Asp Arg Trp Gly Ala Arg 115 120 125

Cys Glu Phe Pro Cys Ala Cys Gly Pro His Gly Arg Cys Asp Pro Ala 130 135 140

Thr Gly Val Cys His Cys Glu Pro Gly Trp Trp Ser Ser Thr Cys Arg 145 150 155 160

Arg Pro Cys Gln Cys Asn Thr Ala Ala Ala Arg Cys Glu Gln Ala Thr 165 170 175

Gly Ala Cys Val Cys Lys Pro Gly Trp Trp Gly Arg Arg Cys Ser Phe 180 185 190

Arg Cys Asn Cys His Gly Ser Pro Cys Glu Gln Asp Ser Gly Arg Cys 195 200 205

Ala Cys Arg Pro Gly Trp Trp Gly Pro Glu Cys Gl
n Gln Gln Cys Glu 210 215 220

Cys Val Arg Gly Arg Cys Ser Ala Ala Ser Gly Glu Cys Thr Cys Pro 225 230 235 240

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His Gly Val Gln Cys Ala His Ser Cys Gly Arg Cys Lys His As
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Pro Cys Ser Pro Asp Thr Gly Ser Cys Glu Ser Cys Glu Pro Gly Trp 275 280 285

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Ser Cys Glu Gln Gln Cys Pro His Cys Arg His Gly Glu Ala Cys Glu 305 310 315 320

Pro Asp Thr Gly His Cys Gln Arg Cys Asp Pro Gly Trp Leu Gly Pro 325 330 335

Arg Cys Glu Asp Pro Cys Pro Thr Gly Thr Phe Gly Glu Asp Cys Gly

340 345 350

Ser Thr Cys Pro Thr Cys Val Gln Gly Ser Cys Asp Thr Val Thr Gly 355 360 365

Asp Cys Val Cys Ser Ala Gly Tyr Trp Gly Pro Ser Cys Asn Ala Ser 370 375 380

Cys Pro Ala Gly Phe His Gly Asn Asn Cys Ser Val Pro Cys Glu Cys 385 390 395 400

Pro Glu Gly Leu Cys His Pro Val Ser Gly Ser Cys Gln Pro Gly Ser 405 410 415

Gly Ser Arg Asp Thr Ala Leu Ile Val Gly Ser Leu Val Pro Leu Leu 420 425 430

Leu Leu Phe Leu Gly Leu Ala Cys Cys Ala Cys Cys Cys Trp Ala Pro 435 440 445

Arg Met Lys Leu Gln Val Trp Gly Thr Leu Thr Ser Leu Gly Ser Thr 465 470 475 480

Leu Pro Cys Arg Ser Leu Ser Ser His Lys Leu Pro Trp Val Thr Val
485 490 495

Ser His His Asp Pro Glu Val Pro Phe Asn His Ser Phe Ile Glu Pro 500 505 510

Pro Ser Ala Gly Trp Ala Thr Asp Asp Ser Phe Ser Ser Asp Pro Glu 515 520

Ser Gly Glu Ala Asp Glu Val Pro Ala Tyr Cys Val Pro Pro Gln Glu 530 535 540

Gly Met Val Pro Val Ala Gln Ala Gly Ser Ser Glu Ala Ser Leu Ala 545 550 555 560

Ala Gly Ala Phe Pro Pro Pro Glu Asp Ala Ser Thr Pro Phe Ala Ile 565 570 575

Pro Arg Thr Ser Ser Leu Ala Arg Ala Lys Arg Pro Ser Val Ser Phe

580 585 590

Ala Glu Gly Thr Lys Phe Ala Pro Gln Ser Arg Arg Ser Ser Gly Glu 595 600 605

Leu Ser Ser Pro Leu Arg Lys Pro Lys Arg Leu Ser Arg Gly Ala Gln 610 615 620

Ser Gly Pro Glu Gly Arg Glu Ala Glu Glu Ser Thr Gly Pro Asp Glu 625 630 635 640

Ala Glu Ala Pro Glu Ser Phe Pro Ala Ala Ala Ser Pro Gly Asp Ser 645 650 655

Ala Thr Gly His Arg Arg Pro Pro Leu Gly Ser Arg Thr Val Ala Glu $660 \hspace{1.5cm} 665 \hspace{1.5cm} 670$

His Val Glu Ala Ile Glu Gly Ser Val Gln Glu Ser Ser Gly Pro Val 675 680 685

Thr Thr Ile Tyr Met Leu Ala Gly Lys Pro Arg Gly Ser Glu Gly Pro 690 695 700

Val Arg Ser Val Phe Arg His Phe Gly Ser Phe Gln Lys Gly Gln Ala 705 710 715 720

Glu Ala Lys Val Lys Arg Ala Ile Pro Lys Pro Pro Arg Gln Ala Leu 725 730 735

Asn Arg Lys Lys Gly Ser Pro Gly Leu Ala Ser Gly Ser Val Gly Gln 740 745 750

Ser Pro Asn Ser Ala Pro Lys Ala Gly Leu Pro Gly Ala Thr Gly Pro 755 760 765

Met Ala Val Arg Pro Glu Glu Ala Val Arg Gly Leu Gly Ala Gly Thr 770 775 780

Glu Ser Ser Arg Arg Ala Gln Glu Pro Val Ser Gly Cys Gly Ser Pro 785 790 795 800

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cga c Arg A 5	egt Arg 50	tca Ser	gtg Val	tct Ser	ttc Phe	cct Pro 55	att Ile	tat Tyr	tgt Cys	gca Ala	agc Ser 60	tta Leu	gca Ala	tat Tyr	gca Ala	192
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ctg g Leu G																288
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	aga Arg															768	
	gat Asp															816	
	gcg Ala															864	
	tac Tyr 290															912	
	gaa Glu															960	
	caa Gln															1008	
~~~	cca	ata	000	~~~	+	~+ n	o at	~ <del>+</del> +								1056	
	Pro										att Ile					1056	
Gly		Val	Pro 340 att	Ala	Ser	Val ctt	Thr	Val 345 tta	Thr	Gly	Ile	Asn	Val 350 cca	Thr	Gln aag	1104	
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gct Ala ggg Gly 385 cca Pro tat Tyr	cca Pro gga Gly 370 tca Ser att	Val aca Thr 355 aaa Lys aaa Lys tat Tyr aac Asn	Pro 340 att Ile ttc Phe att Ile aat Asn aca Thr 420 tta	ggc Gly act Thr acc Thr gga Gly 405 aaa Lys	gaa Glu cca Pro att Ile 390 aat Asn aat Asn	val ctt Leu ttg Leu 375 aca Thr att Ile tac Tyr	Caa Gln 360 ttc Phe tac Tyr acg Thr gcc Ala	Val 345 tta Leu tat Tyr agt Ser tct Ser tgg Trp 425 aga	ttc Phe tca Ser gat Asp gct Ala 410 gcc Ala	act Thr tct Ser gca Ala 395 tta Leu tgt Cys	ctt Leu tca ser 380 aca Thr aat Asn agt ser	att Ile 365 tct Ser aac Asn gac Asp att Ile	Val 350 cca Pro ttt Phe cca Pro tgc Cys tcc ser 430 gat	gaa Glu gat Asp gca Ala aaa Lys 415 gaa Glu aga	Gln aag Lys gac Asp aat Asn 400 tat Tyr atc Ile	1104 1152 1200 1248	

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gat Asp	tgc a Cys I	ct c	Slu I	ccg ( Pro 1 965	gct t Ala C	gt gt Cys Va	a aa al As	t gg n G1 97	y Gl	t gt y Va	t cgg l Arg	gat Asp	ggg Gly 975	Ser	2928
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gag Glu	Ser A	agg ( Arg ( 995	caa a Gln <i>l</i>	aat Asn 1	ctg c Leu E	ro I	a a Le <i>P</i>	ga a rg I	aa t ys I	ta t eu S	er Th	a c r P 05		at tac sn Tyr	3024
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	act Thr 1040					tct Ser 1045					gct Ala 1050		agc Ser		3159
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atc Ile	atg Met 1115	Leu	acc Thr	aat Asn	aat Asn	ttg Leu 1120	Asn	gga Gly	agt Ser	tta Leu	caa Gln 1125	ttg Leu	aac Asn	aat Asn	3384
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	act Thr 1205	gca Ala	gca Ala	ggc Gly	tta Leu	gaa Glu 1210	gtt Val	ttc Phe	aac Asn	aag Lys	agc Ser 1215	gcg Ala	att Ile	gta Val	36	54
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Glu Thr Ala Gln Thr Glu Gln Ser Ser Asp Tyr Gly Thr Phe Leu Thr 165 170 175

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Val Ile Val Ala Asp Val Asn Ala Leu Ser Thr Ser Ser Leu Ser Asn 545 550 555 560

Thr Val Pro Gly Ile Thr Gly Val Pro Ile Thr Val Ile Leu Tyr Gly 565 570 575

Asp Gly Asn Ala Pro Arg Ile Val Gln Ser Thr Thr Asn Ser Glu His

580 585 590

Leu Ser Glu Val Leu Gly Ala Pro Thr Gly Thr Thr Thr Ser Pro Ser 595 600 605

Ser Thr Ala Ser Pro Ala Pro Ser Ala Ser Asp Met Tyr Lys Ala Leu 610 620

Asn Leu Ala Leu Asp Asn Gln Leu Thr Asp Arg Ala Phe Val Ile Val 625 630 635 640

Tyr Thr Ser Asn Ser Asn Val Thr Ile Asp Pro Asp Phe Leu Ile Arg 645 650 655

Leu Ala Val Arg Arg Ala Glu Val Arg Val Leu Ser Thr Ser Gly Thr 660 665 670

Glu Ser Gln Asn Ser Ile Thr Leu Ser Leu Val Gly Asn Gly Ile Pro $675 \hspace{1.5cm} 680 \hspace{1.5cm} 685$ 

Ile Ala Ile Asn Gly Ala Lys Asp Phe Glu Asn Tyr Leu Thr Tyr Tyr 690 695 700

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Val Tyr Asn Gln Gln Asn Thr Leu Asn Ile Tyr Ser Ala Pro Phe Ser 835 840 845

Gln Leu Pro Lys Thr Asp Pro Ser Asn Phe Pro Tyr Leu Thr Leu Gly 850 860

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Asp Val Cys Ser Gln Ser Thr Thr Ile Gly Tyr Gln Ala Gln Leu Thr 900 905 910

Thr Arg Thr Ser Thr Leu Thr Asn Arg Ile Gln Gln Phe Val Leu Thr 915 920 925

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Gly Ser Pro Met Ala Thr Cys Ala Cys Asn Ala Asp Phe Gly Gly Ile 945 950 955 960

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Val Cys Arg Cys Pro Val Arg Phe Tyr Gly Leu Arg Cys Glu Glu Cys 980 985 990

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Leu Gln Ser Asn Ile Lys Ile Ala Ile Asp Arg Ser Asn Ser Ser

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Asn Pro Glu Val Asn Thr Ile Met Asn Asn Phe Leu Val Asn Phe 1295 1300 1305

Arg Leu Gly Phe Asn Tyr Ile Glu Asn Asp Asp Asn Arg Cys Gly 1310 1320

Asp Lys Leu Pro Glu Phe Tyr Thr Gly Arg Thr Ala Leu Thr Ala 1325 1330 1335

Phe Pro Tyr Tyr Asp Ala Ile Ala Lys Ile Ala Thr Phe Cys Pro 1340 1345 1350

Leu Gln Pro Asn Phe Thr Asp Lys Tyr Val Lys Val Ile Ser Lys 1370 1375 1380

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Asp Gln Glu Cys Thr Ile Pro Ile Cys Glu Gly Pro Asp Ala Cys Gln 50 60

Lys Asp Glu Val Cys Val Lys Pro Gly Leu Cys Arg Cys Lys Pro Gly 65 70 75 80

Phe Phe Gly Ala His Cys Ser Ser Arg Cys Pro Gly Gln Tyr Trp Gly 85 90 95

Pro Asp Cys Arg Glu Ser Cys Pro Cys His Pro His Gly Gln Cys Glu 100 105 110

Pro Ala Thr Gly Ala Cys Gln Cys Gln Ala Asp Arg Trp Gly Ala Arg 115 120 125

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Ala Cys Arg Pro Gly Trp Trp Gly Pro Glu Cys Gln Gln Gln Cys Glu 210 215 220

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- Pro Cys Ser Pro Asp Thr Gly Ser Cys Glu Ser Cys Glu Pro Gly Trp 275 280 285
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- Glu Ala Lys Val Lys Arg Ala Ile Pro Lys Pro Pro Arg Gln Ala Leu 725 730 735

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- Lys Ser Lys Glu Gln Glu Glu Leu Leu Gln Met Ile Gln Asn Leu Gln 165 170 175
- Glu Ala Leu Gln Arg Ala Ala Asn Ser Ser Glu Glu Ser Gln Arg Glu 180 185 190
- Leu Lys Gly Lys Ile Asp Thr Leu Thr Leu Lys Leu Asn Glu Lys Ser 195 200 205
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- Lys Asn Arg Gln Thr Cys Gln Ser Leu Gly Gly Gln Leu Leu Gln Ile 260 265 270
- Asn Gly Ala Asp Asp Leu Thr Phe Ile Leu Gln Ala Ile Ser His Thr 275 280 285
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